Salivary Glucosyltransferase B as a Possible Marker for Caries Activity

A.M. Vacca Smitha K.M. Scott-Anneb M.T. Whelehanbc R.J. Berkowitzc
C. Fendd W.H. Bowend

aDepartment of Microbiology and Immunology, bCenter for Oral Biology, cEastman Department of Dentistry and dDepartment of Biostatistics and Computational Biology, University of Rochester, Rochester, N.Y., USA

Key Words
Dental caries · Glucosyltransferase B · Saliva · Streptococcus mutans

Abstract
Bacteria-derived glucosyltransferases (Gtf) (EC 2.4.1.5), through synthesizing glucan polymers from sucrose and starch hydrolysates, play an essential role in the etiology and pathogenesis of caries. We attempted to correlate the levels of Gtf in whole saliva with the prevalence of carious lesions in young children. We examined saliva from children who were either free of overt carious lesions, or had severe early childhood caries (mean dmfs = 18.72 ± 9.0 SD), for Gtf by direct enzyme assay. The levels of GtfB, GtfC and GtfD from Streptococcus mutans in the saliva using monoclonal/specific antibodies in an enzyme-linked immunosorbent assay were determined. Multiple logistic regression analyses with model selection showed that GtfB levels correlated with dmfs values of the subjects (p = 0.006). There was no correlation between total Gtf activity as measured by direct enzyme assay and dmfs values. There was a strong correlation between mutans streptococci populations in saliva and caries activity. Collectively, these data show that GtfB levels in saliva correlate strongly with presence of clinical caries and with number of carious lesions in young children. It is also possible to measure different GtfS, separately, in whole saliva. These observations may have important clinical implications, may lead to development of a chair side caries activity test and support the importance of GtfB in the pathogenesis of dental caries.

KARGER
© 2007 S. Karger AG, Basel

William H. Bowen, BDS, PhD
University of Rochester, Center for Oral Biology
601 Elmwood Avenue, Box 611
Rochester, NY 14642 (USA)
Tel. +1 585 275 0772, Fax +1 585 276 0190, E-Mail William_Bowen@urmc.rochester.edu

Accessible online at:
www.karger.com/cre
S. mutans, S. sobrinus and S. sanguinus [Hamada and Slade, 1980; Vacca Smith and Bowen, 2000].

Clearly GtfS play a critical role in the pathogenesis of caries and furthermore, the enzymes are present in whole saliva. We therefore hypothesized that the amount of GtfS collectively or separately could be correlated with caries saliva. We therefore hypothesized that the amount of GtfB could have predictive value.

Materials and Methods

Antibodies

Polyclonal antisera to Gtfs C and D were raised in rabbits to purified GtfC or GtfD [Venkitaraman et al., 1995; Wunder and Bowen, 2000] and were made monospecific for their target Gtfs by exposing these antisera to hydroxyapatite beads coated with purified nontarget Gtfs, thereby removing antibodies cross-reactive to nontarget Gtf enzyme (unpublished data). Monospecificity of antibodies in the antisera was determined by ELISA assay (see method below) using purified GtfB, GtfC and GtfD as target antigens. ELISA and Western blot assays were also performed to verify that the antibodies did not cross-react with salivary constituents.

Monoclonal antibody to GtfB was prepared by fusion of spleen cells from mice immunized with purified GtfB [Venkitaraman et al., 1995] with myeloma cell line (ATCC/TIB-9) and cloning by limiting dilution using the techniques described by Ivanyi and Davies [1980]. The specificity of monoclonals in hybridoma cell line culture supernatant fluids was determined by ELISA assay (see method below) using purified GtfB, GtfC and GtfD as target antigens, and the antibody was found specific for GtfB.

All animal manipulations were performed in accordance with guidelines established by the University of Rochester Committee on Animal Research.

Dental Examination and Saliva Collection

Whole saliva was collected from 50 children, 25 of whom were diagnosed as having ECC and 25 were clinically caries-free, at the time of their dental examination. All of the ECC subjects were recipients of New York State Medicaid, indicating that they were from families of low economic status. The caries-free subjects had private insurance, showing that they were middle-income families. We chose to carry out our study in children with ECC because we wanted to ensure that our subjects were indeed caries-active. Human subject recruitment, consent and saliva collection were performed in accordance with a protocol approved by the University of Rochester Institutional Review Board. Every subject was given a unique identifier and information about the subjects was kept confidential in accordance with HIPAA regulations. Study subjects were recruited from the patient population of the Division of Pediatric Dentistry, Eastman Dental Center, and University of Rochester. The criteria for establishing a diagnosis of severe ECC were carious lesions affecting at least 2 of the 4 maxillary primary incisors and 2 of the 4 buccal segments.

Caries status was evaluated by 2 trained and calibrated clinical examiners (M.T.W. and R.J.B.) at the time of entry of the subjects into the study. The examinations were done with the aid of a dental mirror and explorer after the teeth had been dried with compressed air. The ECC subjects had their dental exam performed in hospital operating rooms after general anesthesia had been induced. The caries-free subjects had their examination performed during a visit to the pediatric outpatient clinic while they sat in a dental chair. The examiners were recalibrated every 3 months during the study period. No opportunity was provided for performing repeated evaluations on the same subject study by the same examiner and thus no quantitative assessment of intraexaminer reliability was calculated. A surface was declared as having carious lesions per the criteria of Radike [1972]. White spot lesions were included in the caries scoring, they were not penetrated with an explorer.

An unstimulated whole saliva sample was obtained from each subject. The sample was obtained through a disposable saliva ejector attached to a 15-ml sterile centrifuge tube, which in turn was attached to a vacuum pump [Leverett et al., 1993a]. Approximately 2 ml of saliva was collected from each subject. Group A subjects had their saliva sample taken before their oral rehabilitation under general anesthesia and accordingly had nothing to eat or drink for at least 8 h prior to collection. The parents of group B subjects were instructed to give their children no food or beverage for 2 h prior to saliva collection.

After collection, the saliva was immediately transported on ice to the laboratory and assayed within 1 h of collection.

Protein Quantitation

The saliva was clarified by centrifugation, and the amount of protein in the clarified saliva was determined by ninhydrin analyses after wet washing [Moore and Stein, 1948] using glycine as a standard.

Gtf ELISA

We determined specific Gtf (B, C, D) present in the saliva samples using an ELISA assay by employing a kit assay obtained from Kirkegaard and Perry Laboratories, Gaithersburg, Md., USA [Voller et al., 1979; Clark and Engval, 1980]. Clarified saliva was mixed in a 1:1 ratio with the coating buffer supplied in the kit and was coated onto 96-well plates. After washing, antibodies to Gtf B, C and D were applied. All of the buffers and reagents, which were supplied in the kit, were prepared according to the manufacturer’s instructions, and the assays were performed according to the methods outlined in the manufacturer’s information. Controls included wells which did not contain saliva (negative) and wells which did not contain saliva but instead contained known concentrations (1 mg of protein) of purified GtfB, GtfC or GtfD (purified by our previous methods [Venkitaraman et al., 1995; Vacca Smith et al., 1996a]). After development, the intensity of the color in the wells, which correlated to the amount of purified Gtf present, was read in an ELISA reader (Bio-Rad, Hercules, Calif., USA), and the values obtained from experimental and positive control samples were each divided by the negative control values. The results from the calculations were then termed ‘Absorption Index value’. An Absorption Index value ≤ 1 would indicate that the experimental and positive control samples were not different from the negative controls, while an Absorption Index value >1 would indicate that the experimental and positive control samples were different from the negative control.
Salivary Glucosyltransferase B as a Caries Activity Marker

**Gtf Activity**

Gtf activity was determined in all salivary samples by means of direct enzyme assay [Schilling and Bowen, 1988]. A measured volume of clarified saliva from each subject was exposed to 14C-glucosyl-sucrose (final concentration = 100 mmol/l sucrose, 40 μmol/l dextran 9,000 in buffered-KCl, pH 6.5) for 4 h at 37 °C, to form glucans. Gtf activity by direct enzyme assay was expressed as micromoles of glucans formed per milliliter saliva and also per microgram salivary protein.

**Microbiological Analyses**

The levels of mutans streptococci (MS) in the saliva of children from both groups were determined by plating a 20-μl portion of the uncentrifuged (diluted 1:6) saliva on selective medium (mitis salivarius agar + bacitracin) [Gold et al., 1973] and calculating the number of colony forming units (CFU) of MS per milliliter saliva.

**Statistical Analyses**

The subjects were divided into 2 groups, those with carious lesions and those who were lesion-free. The 2-sample t-test was used to determine the significance of differences for CFU, GtfB, GtfC values and Gtf activity. The differences between the 2 groups were also examined using the Wilcoxon rank-sum test. The significance value was set at 0.05. Multiple logistic regression (with model selection) with CFU, GtfB, GtfC, GtfD, Gtf Act and protein as covariates was used to study the effects of these variables on the probability of the occurrence of number of carious lesions. Data analyses were performed using SAS 9.2 (SAS Institute Inc., Cary, N.C., USA). Receiver-operating characteristic plots were used to determine the sensitivity and specificity with MedCalc version 9201 (MedCalc Software, Mariakerke, Belgium).

**Results**

Data on dmfs were determined on 50 children, 25 of whom were caries-free and 25 of whom had severe ECC. The mean dmfs of the children with severe ECC was 18.72 ± 9.0 SD. The ECC group consisted of 15 males and 10 females ranging in age from 38 to 70 months (mean age = 55 months) and comprised 10 African-Americans, 11 Caucasians, 2 Asians, 1 Hispanic and 1 mixed racial. The caries-free group subjects consisted of 15 males and 10 females ranging in age from 42 to 70 months (mean age = 48 months) and comprised 4 African-Americans, 19 Caucasians, 1 Asian and 1 Hispanic.

Table 1 summarizes the results comparing values in saliva from children who were caries-active compared with those who were caries-free. The Absorption Index value for GtfB and the MS concentration were significantly different between the 2 groups. No other values showed a significant difference.

The Gtf Absorption Index correlated significantly with dmfs (p value = 0.006), with an odds ratio (OR) of 14.1 (95% CI = 2.1–94.9). Neither GtfC nor GtfD values, as measured by ELISA, correlated with dmfs. The sensi-

<table>
<thead>
<tr>
<th>Measure</th>
<th>Sensitivity, %</th>
<th>Specificity, %</th>
<th>Cutoff</th>
<th>Area under the curve</th>
</tr>
</thead>
<tbody>
<tr>
<td>GtfB Absorption Index</td>
<td>84</td>
<td>68</td>
<td>1.5</td>
<td>0.782</td>
</tr>
<tr>
<td>GtfC Absorption Index</td>
<td>80</td>
<td>56</td>
<td>1.06</td>
<td>0.634</td>
</tr>
<tr>
<td>GtfD Absorption Index</td>
<td>88</td>
<td>24</td>
<td>2.16</td>
<td>0.491</td>
</tr>
<tr>
<td>Gtf activity (per milliliter saliva)</td>
<td>75</td>
<td>52</td>
<td>0.89</td>
<td>0.571</td>
</tr>
<tr>
<td>Gtf activity (per microgram salivary protein)</td>
<td>79</td>
<td>56</td>
<td>1.10</td>
<td>0.570</td>
</tr>
<tr>
<td>MS, CFU/ml</td>
<td>75</td>
<td>92</td>
<td>3.3 × 10^4</td>
<td>0.853</td>
</tr>
</tbody>
</table>

1 Micromoles of glucose incorporated into glucan/milliliter of saliva.

2 Micromoles of glucose incorporated into glucan/microgram of salivary protein; original value multiplied by 1,000.
tivity and specificity of GtfB as an indicator of caries activity, determined using receiver-operating characteristic plots, were 84 and 68%, respectively (table 2).

Gtf activity as measured by direct enzyme assay failed to show any significant correlation with caries experience (table 1), whether expressed in terms of milliliters saliva or of micrograms protein.

A strong correlation was observed between the number of mutans and caries experience (table 1). The logistic regression of dmfs on CFU showed significant effect (p = 0.01; OR = 1.04, 95% CI = 1.01–1.08). The sensitivity and specificity of MS populations in detecting caries activity are shown in table 2. A correlation between Gtf values in saliva and populations of MS was not detected.

A correlation was not observed between protein concentration of the saliva and caries experience.

**Discussion**

Our study revealed a strong correlation between the level of GtfB, as determined by ELISA using monoclonal antibodies, with the number of clinical lesions of our pediatric subjects. The ability to measure the level of a proven virulence factor and correlate it with clinical caries activity represents a step forward toward using the assay to aid in diagnosis of caries before overt lesions are present.

The failure to find a correlation between total Gtf activity and dmfs scores was surprising. It is possible that a significant amount of activity was removed when the saliva was clarified, through adsorption of enzymes to surfaces of bacteria [Vacca-Smith et al., 1996b]. The activity assay measures both soluble and insoluble glucan formation [Schilling and Bowen, 1992], whereas the ELISA as used here assesses individual enzymes [Yamashita et al., 1993]. Further, it is possible that the assay measures Gtf activity from noncariogenic organisms and clearly does not distinguish different types of Gtf activity.

Our data confirmed previously reported correlations between salivary MS populations and dmfs [Alaluusua and Renkonen, 1983; Köhler et al., 1988; Berkowitz, 1996; Mattos-Graner et al., 2000]. The correlation between salivary GtfB and dmfs appears to be stronger, based on OR, than that between MS and dmfs. In an attempt to determine whether multiple salivary MS counts could provide better predictive power than a single assessment, Petti and Hausen [2000] examined 304 initially caries-free 6-year-olds. The predictive power of a single test had a sensitivity of 29.1% and a specificity of 95.4%. Using multiple MS tests and comparing 0–1 positive tests with 2–3 positive tests, the sensitivity increased to 31.8% and the specificity to 97.6%. Several authors [Grindefjord et al., 1995; van Palenstein et al., 2001; Holbrook et al., 1993] have used combinations of predictors including salivary MS to identify children at high risk to develop caries. Salivary MS counts contributed very little to identifying the at-risk groups. Using the Strip mutans chair side method to assess MS, the predictive ability varied in sensitivity 65% and specificity 86% in a low fluoride group to 40% (sensitivity) to 91% (specificity) in an optimum fluoride group. Our data are not strictly comparable because we carried out a cross-sectional study in young populations with contrasting levels of caries activity. Although the laboratory assessments were done blindly, cross-sectional studies have limitations.

We believe that using GtfB as a potential marker for caries activity has considerable advantages over cultural methods and other approaches such as PCR. GtfB is a proven virulence property of S. mutans. Therefore, the test proposed here includes an agent directly involved in the pathogenesis of the disease. The test as envisaged could be used chair side and provide results within minutes. It is difficult to imagine primer and PCR mutans counts being used chair side, however carried out, as they require days to develop and need some laboratory equipment. In addition, as Bowden [1997] pointed out, cultural methods do not differentiate among clonal types which may or may not be associated with virulence. This observation may also in part account for the relatively poor performance of MS counts as predictors of caries development. It may also explain why a correlation between MS counts and Gtf was not observed in this study (data not shown).

Data from a diverse range of different tests show a correlation with dmfs or DMFT in large populations, especially adults. For example, some tests determine the salivary and plaque populations of cariogenic micro-organisms such as lactobacilli and streptococci [Rogosa et al., 1951; Duchin and van Houte, 1978; Köhler and Bratthall, 1979; Beighton, 1991; Eisenberg et al., 1991; Leverett et al., 1993a, b]. Other methods have been used to identify aciduric and acidogenic organisms in saliva or in plaque, such as the Swab test and the Snyder test [Snyder, 1951; Grainger et al., 1965]. These tests are based on colorimetric measure of pH changes in culture media inoculated with either saliva or plaque samples. They appear to have little predictive value as applied to individuals and generally cannot be conducted at chair side.

An ideal test for caries activity is one that could determine caries activity prior to the onset of lesions. The best
predictor of future caries experience thus far involves assessing the presence of carious lesions already present on tooth surfaces [Grainger and Nikiforuk, 1960; Stamm et al., 1993; Hausen 1997; Powell 1998; Messer, 2000], which from a clinical perspective is rather late.

Our data represent an essential first step in developing a simple and reliable caries activity and predictive test. The strong correlation between lesions and GtfB is certainly encouraging. A problem with our study and indeed all cross-sectional studies is that one has to assume that all clinical caries-free subjects are caries-inactive at the time of sampling. We have a strong suspicion that this was not necessarily so because 2 children who were considered caries-free but displayed high Gtf levels were found to have carious lesions 3–6 months after initial examination. Clearly, this difficulty could be obviated by a longitudinal study in similar populations. The sum of sensitivity and specificity should reach a value of 140 at least for high-risk prediction [Stamm et al., 1993]. Our data show that the GtfB and CFU/milliliter MS met this value (152 vs. 167). Clearly, the true clinical value of our approach remains to be determined until the completion of a longitudinal study.

ECC constitutes a major public health problem and affects those who are least able to bear either the financial or health burden. Children in underserved areas visit a physician more readily than a dentist. If a simple test is developed, such as described here, it could be used by the physician to screen the children for caries risk during ‘well child’ visits and then make appropriate referrals of those identified as high risk to a dentist for the institution of preventive procedure, perhaps reducing the incidence of dental caries, obviating the need for extensive restorative care.

Acknowledgments

This study was supported by USPHS R21DE015564. We are grateful to Dr. John Daiss for discussion of ROC analysis.

References


