

Novel and conventional assays in determining abundance of *Streptococcus mutans* in saliva

Abstract

Objective: This study aimed to comparatively evaluate the validity of a novel immunoassay and a conventional assay in detecting salivary *Streptococcus mutans* (*S. mutans*) in a pediatric cohort.

Methods: A total of 190 children aged 3-4 years were recruited from three kindergartens representing low, moderate, and high socio-economic groups in Hong Kong. The abundance of *S. mutans* in the saliva samples was analyzed with three assay systems viz. a conventional culture-based assay (Dentocult SM), a novel immunoassay system (Saliva-Check *mutans*) based on monoclonal antibody technology, and a Taqman real-time PCR assay taken as a gold standard.

Results: The immunoassay accurately categorized salivary samples into two groups with high ($\geq 5 \times 10^5$ CFU/ml) and low ($< 5 \times 10^5$ CFU/ml) *S. mutans* levels. The sensitivity/ specificity was 97.6%/90.6%. The conventional culture-based assay reached a reasonably high sensitivity/specificity (92.8%/81.3%) in identifying children with moderate ($\geq 10^4$ CFU/ml) *S. mutans* level. However, its sensitivity/specificity in selecting children with high ($\geq 10^5$ CFU/ml) and very high ($> 10^6$ CFU/ml) *S. mutans* levels were not sufficient (78.7%/79.8% and 25.8%/91.8%, respectively).

Conclusions: The monoclonal antibody-based immunoassay, accurately and rapidly determining *S. mutans* abundance in saliva, could be useful for assessing caries risk of patients.

1. Introduction

The use of salivary biomarkers for identifying caries-susceptible individuals is of great interest to dental professionals, while they are striving toward cost-effective caries control and optimized dental treatment planning.¹ Since the central role of *Streptococcus mutans* (*S. mutans*) in the initiation of caries has been well established^{2,3}, abundance of *S. mutans* in saliva is one of the most commonly used microbiological parameters for assessing caries risk.

A variety of test assays are commercially available for measuring the abundance of *S. mutans* in saliva.^{4,5} Of these widely used are the culture-based detection assays. A noted drawback of these culture-based methods is that they are not sufficiently specific to *S. mutans*. Culture-based methods detect not only *S. mutans*, but also other members of the *mutans* Streptococcus group and even some non-Streptococcus species, most of which are not cariogenic.^{4,5} Moreover, conventional culture-based assays normally require 2-4 days to obtain a result as cultures need to be incubated for sufficient time before enumeration.⁴ This appears to be inconvenient for patients, clinicians and public health workers, hampering the wide applications of these products.

Recently, a *S. mutans* detection system Saliva-Check[®] *mutans* immunoassay (GC, Japan) has been introduced. This assay is based on monoclonal antibodies specific for *S. mutans* and the detection depends on antigen-antibody reactions.⁶ Hence, the immunoassay has been designed in such a way that it can offer highly specific and immediate outcome (in 15 minutes). Thus, it appears to be a promising tool for caries risk assessment in the community and clinical settings.

The original Saliva-Check[®] *mutans* system was designed to categorize *S. mutans* abundance into three levels (high, moderate, and low), with the aid of a silver enhancer. A

previous study in a small sample of young adults (n=28) suggested a moderate correlation ($r=0.63$) between Saliva-Check[®] *mutans* readings and the *S. mutans* count, using “PCR-confirmed colony forming unit (CFU)” as a benchmark.⁷ A further comparison showed that the average *S. mutans* count (CFU/ml saliva) in samples classified as with “high” bacterial level was $3.4 \times 10^6 \pm 4.3 \times 10^6$, higher than those for the groups classified as “moderate” ($1.3 \times 10^5 \pm 1.5 \times 10^5$) and “low” ($5.7 \times 10^4 \pm 1.1 \times 10^5$) (both $P < 0.05$). Nevertheless, no significant difference was found between the latter two groups ($P > 0.05$).

Based on these findings, the system has been refined into a two-category assessment outcome (i.e. “high” and “low” denoting *S. mutans* levels above and below 5×10^5 CFU/ml saliva, respectively). However, the validity of this modified immunoassay system has not been evaluated and compared to other detection method in a large-scale epidemiological study and thus is the objective of this study.

2. Method and materials

2.1. Recruitment of participants

An ethical approval for the study was granted by the University of Hong Kong Institutional Review Board (HKU/HA HKW IRB No: UW 08-400). After obtaining parents’ written consents, 190 children aged 3-4 years were recruited from 3 kindergartens located in different districts representing low, moderate, and high socio-economic groups in Hong Kong.

2.2. Determination of *S. mutans* abundance using three methods

The abundance of *S. mutans* in children’s saliva was determined by using a conventional culture-based assay Dentocult[®] SM strip (Orion Diagnostica, Finland), the immunoassay system Saliva-Check[®] *mutans* (GC, Japan), and the Taqman real-time PCR, which served as a

gold standard. During the assessment with each of the three methods, the examiner was blinded to the results obtained from other two methods.

Refraining from foods, drinks, and toothbrushing for at least one hour, each child was instructed to chew paraffin pellet for one minute. The rough surface of the Dentocult[®] SM strip was then pressed on his/her tongue dorsum for 10 times, before being removed between gently closed lips. The Dentocult[®] SM strip was transferred to the microbiological laboratory and incubated in a vial of selective culture broth under 37°C for 48 hours. The density of bacterial colony forming unit (CFU) was rated into four levels (Class 0-3) according to the standard reference chart provided by the manufacturer.

Saliva-Check[®] *mutans* immunoassay system was used to determine the abundance of *S. mutans* in saliva as a chair-side test on the spot. For this purpose, participants were requested to chew paraffin for 3 minutes after the saliva sampling with Dentocult[®] SM strips. Stimulated saliva was collected. A 250- μ l saliva was treated with 50 μ l of Tris-NaOH (Reagent-1) for 30 seconds vigorously, and then mixed with 100 μ l of Tris-citrate (Reagent-2) to neutralize the pH. A 100 μ l sample of the treated saliva was applied to the test device. A red line appearing on the test device after 15 minutes indicated a high *S. mutans* level. If no line was visible, the *S. mutans* level was classified as low.

The remaining portion of the stimulated saliva was transferred to the laboratory and the *S. mutans* level was determined by quantitative species-specific 5' fluorogenic exonuclease (Taqman) real-time PCR (Q-RT-PCR). DNA was extracted from salivary samples as previously described.⁸ DNA extracted from *S. mutans* ATCC cultures (ATCC 35668 and ATCC 700610) was used as the positive control. Q-RT-PCR was performed in ABI PRISM 7900HT sequence detection system using species-specific Taqman probes (Applied Biosciences). Taqman assay

was further validated to quantify the *S. mutans* counts in the range of 10^3 to 10^9 cells. Each PCR was performed in a total volume of 20 μ l consisting 10 μ l of Taqman Gene expression master mix, 1 μ l of forward and reverse primers, 1 μ l of Taqman probe, 1 μ l of template DNA, and appropriate amount of sterilized DNase–RNase-free water. The cycling conditions were 50 °C for 2 minutes, 95 °C for 10 minutes and 40 cycles at 95 °C for 15 seconds, 60 °C for 1 minute.

2.3. Data analysis

The data were analyzed with the Statistical Package for Social Sciences (SPSS, version 17.0). Comparison was made between the mean *S. mutans* counts of two groups defined by Saliva-Check[®] *mutans* as with low and high bacterial load. Non-parametric method (Mann-Whitney test) was applied since the assumption of homogeneity of variance was violated, as shown by the Levene test. For Dentocult[®] SM, means of the four categories (Class 0-3) were compared by using parametric test (post hoc Tukey test), since both the normal distribution and homogeneity of variance were supported. The validity of the immunoassay system and conventional culture-based assay in determining *S. mutans* abundance was evaluated and compared. The main parameters used included sensitivity, specificity, and Youden's index (sensitivity + specificity – 1), which is a single statistic that captures the overall performance of a diagnostic test.⁹ A Youden's index above 0.6 or above 0.8 indicates a “useful” or “good” test, respectively^{10,11}.

3. Results

Table 1 shows the mean number of *S. mutans* [$\log_{10}(\text{CFU/ml saliva})$] confirmed by real-time PCR in each classification of both test assays (Saliva-Check[®] *mutans* and Dentocult[®] SM strip). The immunoassay system Saliva-Check[®] *mutans* indicated 136 (71.6%) and 54 (28.4%) children harbored low and high levels of *S. mutans*, respectively. The mean *S. mutans* count [$\log_{10}(\text{CFU/ml})$] was 3.44 +/- 0.84 for the former group, significantly lower than that of the

latter group, which was 5.97 +/- 0.97 (P<0.001).

The conventional assay Dentocult[®] SM strip classified *S. mutans* counts into four groups in an ascending order from 10^4 CFU/ml to >math>10^6</math> CFU/ml, '0' being the lowest and '3' being the highest. When Taqman real-time PCR assay was used as the gold standard, the *S. mutans* levels [log₁₀(CFU/ml)] of these four categories appeared to be 3.10 +/- 0.56 (Class 0), 4.91 +/- 1.67 (Class 1), 5.20 +/- 1.19 (Class 2), and 5.38 +/- 1.07 (Class 3), respectively. While children in "Class 0" had significantly lower number of *S. mutans*, compared with the rest of children (P<0.001), there was no significant difference among Class 1-3 (P>0.05).

Saliva-Check[®] *mutans* immunoassay accurately categorized the saliva samples into two groups with high ($\geq 5 \times 10^5$ CFU/ml) and low (5×10^5 CFU/ml) *S. mutans* levels (Table 2). The sensitivity/specificity was 97.6%/90.6%. The Youden's Index was 0.882. Dentocult[®] SM strip reached reasonably high sensitivity/specificity (92.8%/81.3%) and Youden's Index (0.741) in selecting children with moderate ($\geq 10^4$ CFU/ml) *S. mutans* level. However, it did not sufficiently identify children with high ($\geq 10^5$ CFU/ml) and very high (>math>10^6</math> CFU/ml) *S. mutans* levels. The sensitivity/specificity were 78.7%/79.8% and 25.8%/91.8%, respectively. Both Youden's indexes were below 0.6. The contingency table (Table 3) compares the classifications by the immunoassay (10^5 CFU/ml as the cut-off point) and culture-based assay (under the best cut-off point of $\geq 10^4$ CFU/ml) against the real-time PCR results.

4. Discussion

Although *S. mutans* is an important parameter for assessing caries risk, its clinical application is limited partly due to the lack of a highly accurate test which can be performed with minimal training and labor. This study, utilizing a relatively large sample and the Taqman real-time PCR technology as the gold standard, comparatively evaluated the validity of a

promising novel immunoassay and a conventional assay. Taqman real-time PCR assay, as an advanced molecular detection technology, is considered one of the best techniques to quantify the bacterial load in clinical samples.¹²

Our results revealed a high validity of the Saliva-Check[®] *mutans* immunoassay system in determining *S. mutans* abundance in saliva. In the present cohort, the system accurately identified about one quarter (28%) of children harboring high level ($\geq 5 \times 10^5$ CFU/ml saliva) of *S. mutans*. The sensitivity and specificity were both above 90%. The Youden's index was 0.882. In light of the current polarized distribution of caries, selecting one quarter of the population with high risk for targeted prevention and intervention is a rational approach for achieving cost-effective caries control and optimized treatment planning.¹³

Comparatively, Dentocult[®] SM strip, with a four-class reading, is meant to generate a more refined, semi-quantitative assessment of bacterial abundance. However, our results indicated a lack of difference in *S. mutans* level among Class 1-3, despite the difference between Class 0 and the rest. Correspondingly, Dentocult[®] SM strip reached a reasonably high sensitivity/specificity (92.8%/81.4%) and Youden's index (0.741) only in identifying children with *S. mutans* level of 10^4 CFU/ml and above (Class 1-3). Our findings suggest that a binary rating system (i.e. $\geq 10^4$ CFU/ml or $<10^4$ CFU/ml) of Dentocult[®] SM strip may better reflect the abundance of *S. mutans*, the main cariogenic bacterial species.

The emergence of molecular detection methods has been one of the landmark revaluations in science over the past century.¹⁴ In the medical arena, these methods offer superior specificity and sensitivity over the conventional culture-based techniques in enumerating microorganism and molecules of health relevance. However, laboratory molecular methods, such as Taqman real-time PCR, are expensive and require specialized skills and equipments. The development

of the immunoassay system for detecting *S. mutans* is in fact an example of introducing the molecular detection method into practical applications at chairside. The immunoassay system is characterized by its rapidity and ease of the testing procedures, which can be completed by a dentist or dental auxiliary. Since no cultivation is needed, the test requires no additional apparatus and minimizes the chance of contaminations. The immunoassay assays can be easily stored under room temperature for at least two years, while the conventional culture-based assays require special storage condition (2-8 °C; protected from light etc), with a shelf life normally no more than 6 months.

Although the immunoassay system provides a practical solution for accurately measuring *S. mutans*, the primary pathogen for caries, it should be borne in mind that *S. mutans* is not the only cariogenic bacterial species related to caries.¹⁵ For further development of the system, immunoassays for detecting other cariogenic species (e.g. *Streptococcus sobrinus*) may be considered. In addition, abundance of cariogenic bacteria should be synthesized with other parameters (diet, fluoride etc) for a better understanding of individual's caries risk.

5. Conclusions

The monoclonal antibody-based immunoassay, accurately and rapidly determining *S. mutans* abundance in saliva, could be useful for assessing caries risk of patients.

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References

1. Tanzer JM, Livingston J, Thompson AM. The microbiology of primary dental caries in humans. *Journal of Dental Education* 2001;**65**(10):1028-37.
2. Keyes PH. The infectious and transmissible nature of dental caries. Findings and implications. *Archives of Oral Biology* 1960;**1**:304-20.
3. Fitzgerald RJ, Fitzgerald DB. The microbiologic status of test animals in relation to caries research. In: Tanzer JM, editor, Animal models in cariology: proceedings of a Symposium and Workshop on Animal Models in Cariology. Washington DC: Information Retrieval Inc. 1981. p.89-95.
4. Tanabe Y, Park JH, Tinanoff N, Turng BF, Lilli H, Minah GE. Comparison of chairside microbiological screening systems and conventional selective media in children with and without visible dental caries. *Pediatric Dentistry*. 2006;**28**(4): 363-8.
5. Nishikawara F, Nomura Y, Imai S, Senda A, Hanada N. Evaluation of cariogenic bacteria. *European Journal of Dentistry* 2007;**1**(1):31-9.
6. Shi W, Jewett A, Hume WR. Rapid and quantitative detection of *Streptococcus mutans* with species-specific monoclonal antibodies. *Hybridoma* 1998;**17**:365–371.
7. Matsumoto Y, Sugihara N, Koseki M, Maki Y. A Rapid and Quantitative Detection System for *Streptococcus mutans* in Saliva Using Monoclonal Antibodies. *Caries Research* 2006;**40**:15–19.
8. Oho T, Yamashita Y, Shimazaki Y, Kushiya M, Koga T. Simple and rapid detection of *Streptococcus mutans* and *Streptococcus sobrinus* in human saliva by polymerase chain reaction. *Oral Microbiology and Immunology* 2000;**15**(4):258-62.
9. Youden WJ. An index for rating diagnostic tests. *Cancer* 1950;**3**:32-35.
10. Schisterman EF, Perkins NJ, Liu A, Bondell H. Optimal cut-point and its corresponding Youden Index to discriminate individuals using pooled blood samples. *Epidemiology* 2005;**16**(1):73-81.

11. Chu CH, Lo EC, You DS. Clinical diagnosis of fissure caries with conventional and laser-induced fluorescence techniques. *Lasers in Medical Science* 2010;**25**(3):355-62.
12. Suzuki N, Yoshida A, Nakano Y. Quantitative analysis of multi-species oral biofilms by TaqMan Real-Time PCR. *Clinical Medical Research* 2005;**3**(3):176-85.
13. Beck JD, Weintraub JA, Disney JA, Graves RC, Stamm JW, Kaste LM et al. University of North Carolina Caries Risk Assessment Study: comparisons of high risk prediction, any risk prediction, and any risk etiologic models. *Community Dentistry and Oral Epidemiology* 1992;**20**(6):313-21.
14. Elliott WH. Biochemistry and Molecular Biology. 4th Edition. Oxford University. 2009.
15. Samaranayake LP. Oral microbiology. In. Essential Microbiology for Dentistry. 3rd Edition. Elsevier. 2006. P. 253-306.