

Salivary Biomarkers in Diagnosis of Dental caries - A Review Article

Review Article

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Abstract

Saliva can be used to study the physiological state of the body, having the potential to be used in early detection and diagnosis of diseases. Saliva is secreted from the salivary glands and has multiple functions, including oral cavity cleaning and protection, antibacterial effects and digestion. With the rapid advancement in salivaomics, saliva is well recognized as a pool of biological markers. Saliva contains various microbes and host biological components that could be used for caries risk assessment. The determination of biomarkers in saliva is becoming an important part of laboratory diagnostics and the prediction of dental caries. Biomarkers in saliva (e.g., enzymes, protein markers, or oxidative stress markers) can be used for activity determination for dental caries prognosis. Saliva is an interesting alternative diagnostic body fluid with several specific advantages over blood. These include non-invasive and easy collection and related possibility to do repeated sampling. This makes saliva very interesting for clinical biochemistry of oral diseases. This review summarizes the latest advancements in saliva-related studies and addresses the potential salivary biomarkers in the early diagnosis of dental caries.

Keywords: Biomarkers; Diagnosis; Dental Caries; Genetic; Salivary Proteins.

Introduction

Biomarker is an objectively measured and evaluated indicator of normal biologic processes, pathogenic processes, or pharmacologic responses to therapeutic intervention. They are entities within the body capable of providing impartial information regarding the current physiologic state of a living organism [1]. Biomarkers exist in a variety of different forms, including antibodies, microbes, DNA, RNA, lipids, metabolites, and proteins. Alterations in their concentration, structure, function, or action can be associated with the onset, progression, or even regression of a particular disorder.

Salivary Biomarkers

The oral cavity is a diverse habitat composed of teeth, gingival sulci, the tongue, hard and soft palates, the buccal mucosa, and tonsils. Each structure is colonized by bacteria and continuously bathed in saliva. Saliva can be used to study the physiological state

of body, having the potential to be used in early detection and diagnosis of diseases [2]. There is an association between elevated caries prevalence and/or incidence among people with a pathologically low saliva flow rate, compromised buffering capacity and early colonization or high titer of mutans streptococci in saliva. Studies have shown that salivary bacteria, may also be used as an indicator of dental caries [3]. There are associations between dental caries and other saliva parameters, such as other cariogenic species (*Lactobacillus* spp., *Streptococcus sanguis* group, *Streptococcus salivarius*, *Actinomyces* spp. and *Candida albicans*), diversity of salivary microbiomes, inorganic and organic constituents (electrolytes, immunoglobulins, other proteins and peptides) and some functional properties [4] (sugar clearance rate, etc).

Sample Collection

The usual salivary collection methods include a draining method using a Proflow Sialometer, a spitting method, a suction method, swab or absorbent methods and the use of salivette. The physi-

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Received: May 04, 2021

Accepted: August 5, 2021

Published: August 14, 2021

Citation: P.S.Subiksha, Raghu Sandhya. Salivary Biomarkers in Diagnosis of Dental Caries - A Review Article. *Int J Dentistry Oral Sci.* 2021;8(8):3729-3733.
doi: <http://dx.doi.org/10.19070/2377-8075-21000764>

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ological pH of saliva ranges from 6.2 to 7.4. Chewing paraffin wax is the most commonly employed saliva stimulating method. The samples are usually collected on ice or any container. The collected samples are immediately centrifuged at 13000 rpm for 5 minutes to remove the insoluble material, and all the procedures are performed at 40 C. The supernatant is removed and placed in Eppendorf tubes that are stored at -800°C [5].

There are many processing units for testing saliva. Most common method performed for testing are enzyme-linked immunosorbent assay (ELISA), polymerase chain reaction (PCR), high resolution mass spectrometry (HRMS), Western blotting technique, newer technology such as fibre-optic-based-detection [6]. The salivary buffer capacity can be measured by Dentobuff method. The saliva Total antioxidant capacity (TAC) measurement can be performed using spectrophotometry by adaptation of 2, 2'-azino-di-(3-ethylbenzthiazoline-6-sulphonate) ABTS assay [7]. Major drawback of ELISA is that it requires more invasive method for collection of samples and trained personnel [8].

Various Salivary Biomarkers

The abundant protein content present in the saliva acts as biomarkers. The various salivary proteins that can be used as biomarkers for dental caries are soluble immunoglobulin A, mucin 1 and 2, cystatin S, statherins, defensins, CD14 and glycosyltransferase. These act as potential markers for dental caries.

Soluble immunoglobulin

Salivary antibodies are the first line defence against antigens present in saliva, epithelial and tooth surface. Salivary immunoglobulins are produced by plasma cells present in stroma of salivary glands, adjacent to salivary ducts and oral mucosa. The main salivary immunoglobulin of whole saliva is secretory immunoglobulin A (sIgA), is constitutively secreted into saliva. It has the ability to inhibit bacterial adhesion to the epithelial cells, and is considered as the first line of defence against bacterial invasion [9]. It acts synergistically with other defence mechanisms, such as the lactoferrins, peroxidase, agglutinins and mucins.

The normal level of sIgA in individuals ranges from 4-30mg/dL. This level is changed by numerous conditions, like malnutrition, obesity, infections, stress, smoking, salivary flow rate, hormonal factors. In elderly persons, a decreased level of sIgA is associated with an increase in root caries and candidiasis.

Based on a study on an Indian population [10], it was observed that the total salivary concentration of sIgA was higher in caries free children than in other groups with active caries. Similarly Omar et al demonstrated that the sIgA levels decreased with the increase of the carious lesions [11]. The group with lower caries experience has a higher level of sIgA than those caries free group. In overview, the levels of sIgA appeared to be inversely proportional to DMFT. Contrary to this, Vitorino et al [12] reported that the IgA was present at higher concentration in caries susceptible group.

All these studies revealed different correlation between the salivary IgA levels and caries. The first two studies found a negative correlation while the last study had a positive correlation, which

associated with higher levels of IgA with higher caries susceptibility.

Mucins 1 and 2

Mucins are glycoproteins produced by submandibular, sublingual, labial and palatal minor salivary glands. They have an important role in concentrations of other antimicrobial proteins in oral mucosa such as lysozymes, IgA and cystatin. Saliva contains two forms of mucins, the high-molecular-weight mucin glycoprotein-1(MG1 or MUC5b) and the low-molecular-weight glycoprotein-2(MG2 or MUC7). These mucins control the process of demineralisation and remineralisation.

In a study on Mexican population, a correlation between the quantity of proteins MG1 and MG2 and DMFT index was observed, where the absence of 6-13% of these mucins was associated with higher DMFT index [13].

Cystatin S and Statherin

Saliva presents seven different cystatins, cystatin A, cystatin B, cystatin C, cystatin D, cystatin S, cystatin SA and cystatin SN[14]. Cystatin S (AA1-8) variant, is a truncated form of cystatin S formed by proteases on the carboxyl side of arginine. Cystatin S can be phosphorylated in five sites. These phosphorylated forms have important function in the regulation of calcium level and in pellicle formation.

Literature search showed that the saliva from caries free group had a higher concentration of Cystatin S, SN1, SN2 and SA-III. The study reported that DMFT had a negative correlation with Salivary Cystatin.[12]

Statherin is a protein with the most important function being the inhibition of precipitation in saturated solutions of calcium. It is the primary regulator of mineralisation in the oral cavity. The salivary levels of statherin and variant cystatin S (AA1-8) have an inverse correlation with occlusal caries [15]. Higher levels of statherin and cystatin S were observed in caries-free individuals. Research report reveals correlation between Cystatin S and Statherin, and bacterial aggregation and adherence. Rudney et al [16] investigated the influence of bacterial aggregation, adherence and killing on the risk of dental caries. The results demonstrated the reduction of caries in groups with high aggregation –adherence. These groups had higher levels of cystatin S and statherin.

Glucosyltransferase B

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. Vacca Smith et al [17] evaluated the salivary glucosyltransferase B as a possible marker for caries activity. They reported that there was a positive relation between mutans streptococci populations in saliva and caries activity. They concluded that GtfB levels in saliva correlate strongly with presence of clinical caries and with number of carious lesions in young children.

Superoxide Dismutase

Hedge et al [18] evaluated the biochemical indicators of dental

caries in saliva with the use of superoxide dismutase (SOD) activity, copper and zinc. They reported that SOD activity as well as copper and zinc levels increased in the caries-active group and showed statistically significant results.

Carbonic anhydrase enzyme

Picco et al [19] evaluated that children with a higher activity of carbonic anhydrase (CA) VI in saliva are more likely to develop dental caries. They reported that the salivary CA VI activity was higher in children with caries. They found a negative correlation between buffering capacity and dental caries. Also, in the caries group they found a positive correlation between the concentration and the activity of CA VI and a negative correlation between BC and CA VI activity. A high activity of CA and a low salivary flow rate were associated with dental caries.

Total antioxidant capacity (TAC)

Mahjoud et al [20] evaluated the comparison of TAC in saliva of children with severe early childhood caries and caries-free children. They compared the TAC levels in the unstimulated whole saliva of children with severe early childhood caries (S-ECC) and caries-free children and concluded that TAC levels and salivary total protein increased in children with S-ECC compared with caries-free children.

Magnetic-bead salivary peptidome

Si et al [21] in their study on magnetic bead-based salivary peptidome profiling analysis for severe early childhood caries proposed a new strategy for screening high-risk populations. Based on a novel method, the salivary protein profiling can be done. They used matrix-assisted laser desorption/ionization time-of-flight mass spectrometry (MALDI-TOF MS) combined with weak cation exchange magnetic beads, and peptide mass fingerprints were created by scanning mass spectrometry signals.

Salivary peroxidase

Lamberts et al [22] evaluated the comparison of salivary peroxidase system components in caries-free and caries-active naval recruit. Based on a single point measurement of saliva no significant correlation between caries incidence and concentrations of salivary peroxidase system was observed. However generation of hypothiocyanite in saliva was associated with a decrease in enzyme activity. There are reports revealing association between smoking and hypothiocyanite. Smokers had significantly elevated thiocyanate and hypothiocyanite compared to nonsmokers. When the salivary peroxidase system is activated, additional supplements of thiocyanate produce increased generation of hypothiocyanite.

Matrix Metalloproteinase (MMP)

Tannura et al [23] evaluated the relation between MMP polymorphisms and dental caries. Matrix metalloproteinases (MMPs) and their tissue inhibitors have been suggested to be involved in the caries process. The MMPs under research are MMP2 (rs243865), MMP9 (rs17576), MMP13 (rs2252070), and TIMP2 (rs7501477) whether they are associated with caries. Genetic variation in MMP13 may contribute to individual difference in caries susceptibility. However allelic and genotype frequencies of the polymor-

phisms in MMP9 were similar in caries affected and caries free individuals.

Nitric oxide (NO) concentration

Bayindir et al [24] evaluated the nitric oxide concentrations in saliva and dental plaque in relation to caries experience and oral hygiene. The result showed that the patients with high DMFT had significantly higher NO concentrations in saliva and plaque than those with low DMFT. Plaque NO concentrations were significantly higher than in saliva. NO production might be a host defence mechanism when dental caries increases or oral hygiene deteriorates.

Genetic

The saliva characteristics are controlled by many factors, including genetic factors. The genetic mechanisms in the aetiology of caries encompass, 4 main groups of genes responsible for (1) the development of enamel, (2) formation and composition of saliva, (3) immunological responses, and (4) carbohydrate metabolism. BMP7, ALOX15, AQP5, TUFT1, KRT75 genes in tooth formation were found to be associated with increased caries susceptibility and a risk factor for caries [25-29]. Haplotypes ACA and GG are associated with high caries susceptibility [30, 31]. These are the genes that are associated with the composition and functions of saliva. DEFB1 and MBL2 are the genes influencing immune response and are found to have a high caries intensity [32, 33]. TAS1R2 and GLUT2 are the genes influencing carbohydrate metabolism and are found to have higher caries intensity [34, 35].

Küchler et al [36] reported that the genes involved in the enamel development are associated with calcium and phosphorus level in saliva. They reported that there were genetic variations in AMELX, AMNB and ESRRB which were associated with the calcium level in saliva and a borderline association was observed in ENAM allele distribution shown with phosphate level in saliva. The antimicrobial peptides human β -defensins (hBDs) are encoded by β -defensin genes (DEFBs) and are possibly involved in caries susceptibility. Lips et al [37] in their study concluded that genetic polymorphism in miRNA202 is involved in hBD1 salivary level as well as caries experience in children.

Gene variants affecting taste preference and glucose transport were recently associated with caries risk. Many clinical studies showed that the difference in sensitivity to the bitter taste of 6-n-propylthiouracil (PR7OP) is a heritable trait and may influence children's caries development. Oter et al [38] in their study evaluated the relation between 6-n-propylthiouracil sensitivity and caries activity in school children. The results concluded that PROP non-tasters were significantly more likely to have high caries risk than PROP tasters. Izakovicova Holla et al analyzed two common polymorphisms in the sweet taste receptor (TAS1R2) and glucose transporter (GLUT2) genes in children with dental caries and healthy controls among the Czech population. They identified that GLUT2 and TAS1R1 polymorphisms influenced the risk of caries. [39, 40]

Our team has numerous highly cited publications on well-designed clinical trials and lab studies on various topics in the past couple of years [41-55]. Our institution is passionate about high

quality evidence based research and has excelled in various fields [56-65]. The current topic was reviewed with an intention to explore the diagnostic tool to be put into research and practice.

Conclusion

Saliva has been analysed for diagnostic purposes. Salivary biomarkers are useful in the diagnosis of variety of diseases. It is non invasive, uncomplicated, diagnostic tool. Dental caries that can be monitored by assaying salivary biomarkers opens a wider view. The dental caries affect the salivary proteome. Consequently saliva appears to be a potential source of biomarkers for dental caries. Based on the analysis of several studies, there are various protein candidate, salivary enzymes, genes which play the role of biomarker for dental caries. More studies on salivary biomarkers may prove greater insight in to various other diseases in human population.

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