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Salivary Glands

New Approaches in Diagnostics and Treatment

Edited by Işıl Adadan Güvenç



SALIVARY GLANDS - NEW APPROACHES IN DIAGNOSTICS AND TREATMENT

Edited by **Işıl Adadan Güvenç**

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Contributors

Iordanis Konstantinidis, Angelos Chatziavramidis, Hisao Hiraba, Michihiro Nakayama, Atsutaka Okizaki, Kaori Nakajima, Koji Takahashi, Alejandro Escobar, Juan Pablo Aitken, Goran Mitulović, Karolina Kaczor-Urbanowicz, Anahita Punj, Işıl Adadan Güvenç

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Meet the editor



Dr. Işıl Adadan Güvenç was born in 1976 in İzmir, Turkey. After graduating from the American Collegiate Institute in İzmir, she received her medical degree at Hacettepe University, Faculty of Medicine, in 2000. She then specialized in otorhinolaryngology head and neck surgery at İzmir Atatürk Research and Training Hospital. Today, she works in Çiğli Regional Training Hospital in İzmir as an associate professor in ENT. She is a fellow of the European Board of ORL-HNS, and has published 15 articles in respectable international journals and 11 articles in national journals. She is married to an architect and has an 8-year-old son.

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Preface

Saliva is a complex fluid that maintains oral health and possesses important functions in lubrication, digestion, taste perception and speech. Moreover, saliva is a potential noninvasive diagnostic fluid, which provides valuable diagnostic information for detecting numerous diseases. Thanks to the advances in diagnostic methods, the number of studies using salivary analysis in the diagnosis of various diseases and conditions has increased tremendously. As evidence from recent research accumulates, salivary diagnostics will become more easily accepted by clinicians and patients.

Salivary Glands - New Approaches in Diagnostics and Treatment is a comprehensive reference, which brings together information on salivary secretion and its disorders, as well as novel diagnostic methods for numerous diseases with the use of salivary secretions, and new techniques in the treatment of salivary diseases. Therefore, this book contains a source of information for a diverse audience, such as researchers of basic science and clinicians of surgical and medical sciences, including dentists, oral biologists, experimental biologists, molecular biologists, oncologists, radiologists, oral and maxillofacial surgeons, and otorhinolaryngologists.

In the first three chapters of the book, a basic understanding of salivary secretion is provided along with a description and management of conditions that decrease or increase the amount of saliva. The next section of the book introduces the reader to salivary diagnostics, a new diagnostic approach that has the potential to reach clinical practice in the near future for the early detection, diagnosis, and monitoring of many diseases, such as cancer, human papillomavirus infections, dental problems, diabetes, heart diseases, Sjögren's syndrome, and even neurological and psychiatric disorders. Afterwards, the proteomics of saliva are described in detail. Next, scintigraphic techniques for the functional diagnosis of salivary diseases and for distinguishing salivary tumors are discussed. The last two chapters describe a combined endoscopic procedure used in recent years in the treatment of sialolithiasis and a novel method, facial vibrotactile stimulation, for the increase of salivation.

I would like to sincerely thank my husband Erkan Güvenç for his patience, my mother Emel Adadan who was always beside me whenever she was needed, and our nanny Ellie Mikrut who took great care of my dearest Kaan for sacrificing their time and supporting me in making this book happen.

Işıl Adadan Güvenç, MD
Cigli Regional Education Hospital
Izmir, Turkey

Salivary Glands and its Diseases

Secretions of Human Salivary Gland

Anahita Punj

Additional information is available at the end of the chapter

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Abstract

The salivary glands play an important role in our body by the virtue of its ability to secrete saliva. Saliva has a role to play in maintaining the health of the oral cavity and for carrying out physiological functions like mastication, taste perception, speech etc. It also acts as a mirror to the systemic status of an individual owing to its ability to act as a diagnostic fluid for detecting a number of conditions and diseases. Saliva is a potential non-invasive diagnostic fluid for detection of a number of biomarkers of disease and health. Advancement in diagnostic methods has helped in identifying biomarkers of disease in saliva. In order to understand and diagnose pathological changes, a thorough understanding of the salivary gland anatomy, physiology and regulation of its secretion is warranted. This chapter aims to provide the basic understanding of the secretions of saliva.

Keywords: saliva, secretion, salivary gland, diagnostic fluid

1. Introduction

Salivary glands are organs which synthesize and secrete their secretions over an epithelial surface via a hollow channel. These glands are present in and around the oral cavity and its secretions play an important role in the physiological processes of the oral cavity [1].

2. Overview of salivary glands

The salivary glands can be classified as major and minor salivary glands. The major salivary glands, located outside the oral cavity include the parotid salivary gland, submandibular/submaxillary salivary gland and sublingual salivary gland. The minor salivary glands are

classified based on their location in the oral cavity as labial/buccal glands, glossopalatine glands, palatine glands, lingual glands which are further classified as anterior lingual (glands of Blandin and Nuhn) and posterior lingual glands (Von Ebner's) [1, 2]. The following diagram (**Figure 1**) shows the anatomical location of major and minor salivary glands. The salivary glands consists of a secretory part and ducts (**Table 1**).

2.1. Parenchymal elements of salivary gland

The salivary glands are made of secretory units called acini, which are made up of acinar cells which could be serous or mucous. The serous cells are pyramidal or triangular in shape while the mucous cells are columnar in shape. The serous cells are occasionally seen capped by structures called demilunes. The acini cells are surrounded by contractile cells called as myoepithelial cells/basket cells, which are responsible for the flow of secretions of saliva by contraction of the cell. The acini of salivary glands are connected to hollow tubular structures which are called salivary ducts. The lining of the duct changes with the type of duct and its location within the salivary gland [1–3]. A description of the ducts observed is given in **Table 2** and the parenchymal elements are shown in **Figure 2**.

2.2. Development of salivary gland

Salivary glands arise from the ectoderm of oral cavity. The minor salivary gland arise from the oral and nasopharyngeal ectoderm. The chronology of the development of salivary gland is mentioned in **Table 3**. Each gland develops at a specific location in the oral cavity by the inward growth of an epithelial bud into the underlying mesenchyme. These epithelial buds then grow and later branch into a system of cords of cells. These get canalized and develop

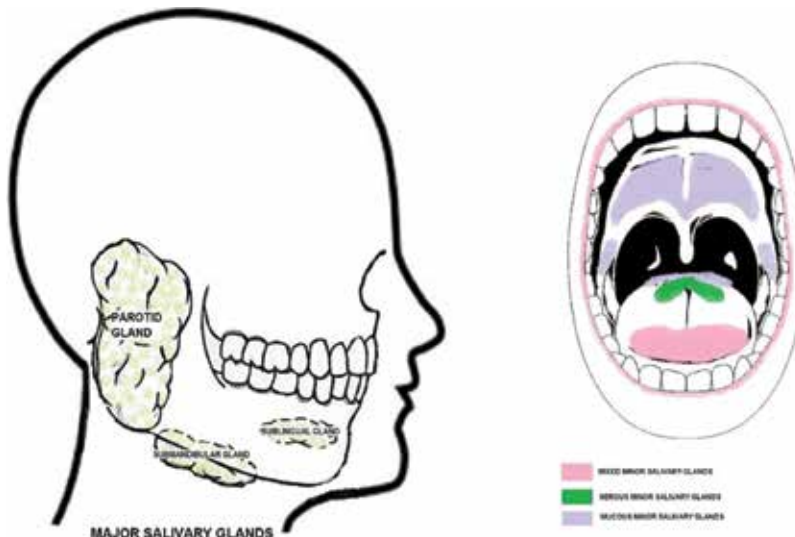


Figure 1. Anatomical locations of major and minor salivary glands.

S. no	Salivary gland	Salivary gland duct	Location of salivary duct orifice
1.	Parotid	Stensen's duct	Opens at papilla in buccal mucosa opposite maxillary second molar
2.	Submandibular	Wharton's duct	Opens at sublingual papillae
3.	Sublingual	Bartholin's duct Duct of Rivinus	Opens with or near submandibular duct Opens independently along sublingual fold
4.	Minor salivary glands	Short ducts	Open directly via short ducts into mouth

Table 1. Location and names of salivary gland ducts.

S. no	Duct	Description	Epithelium
1.	Intercalated duct	Connect the terminal secretory unit with the next system of ducts	Single layer of low cuboidal cells
2.	Striated duct	Intercalated ducts drain into striated duct	Tall columnar epithelial cells with centrally placed nucleus. Cells are partitioned by deep sheet like foldings of membrane, which appear as striations under light microscope
3.	Interlobular duct/ excretory duct	Formed by joining of striated ducts	Pseudo-stratified columnar epithelial cells with outer connective tissue adventitia

Table 2. Description of salivary ducts.

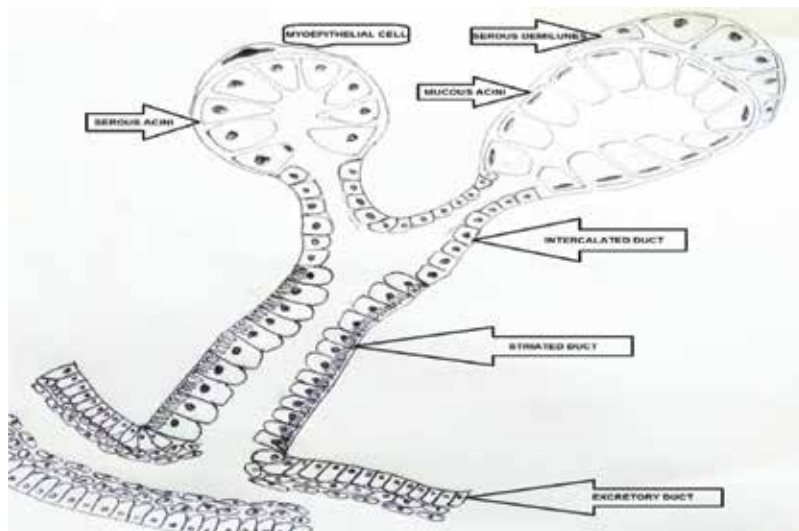


Figure 2. Architecture of salivary ducts and acini.

S. no	Time	Event
1.	Sixth week	Development of the primordia of parotid and submandibular salivary gland
2.	Seventh to eighth week	Development of sublingual salivary gland
3.	Third month	Development of minor salivary gland

Table 3. Chronology of salivary gland development.

a lumen by the action of microfilaments at apical areas of cell to become ducts. The secretory part develops later by repeated branching and budding of finer cell cords, to form pregland cells which give rise to acini [1].

3. Secretions of salivary gland

The salivary glands are primarily involved in secretion of saliva. There are other substances also which are secreted by the salivary glands which are found in saliva. They can secrete proteins in large amounts apically or basolaterally to the saliva [1–4].

3.1. Saliva

Saliva is a complex physiological fluid of the oral cavity which coats the teeth and oral mucosa. It contains a myriad of components like enzymes, mucinous substances, antibacterial components etc. Saliva functions to maintain the oral cavity in the physiological state owing to its lubricating, buffering, antibacterial and immune properties by acting as a physiological barrier to infections. The saliva is a mixed fluid, as it is composed of saliva secreted by both major and minor salivary glands which are both serous and mucous in nature [2–4].

3.1.1. Composition of saliva

Saliva is composed of 99% water and 1% of components such as inorganic ions like sodium, potassium, chloride, bicarbonate, calcium, magnesium, fluoride, thiocyanate, hydrogen phosphate etc. It contains proline-rich proteins, histatins, cystatin, defensins. Kallikrein, cathelicidin-LL37, lactoferrin and enzymes such as amylase (ptyalin), peroxidase, lysozyme, etc. Immunoglobulins A, G and M, glucose, amino acids, urea, uric acid, lipid molecules and blood group antigens, epidermal growth factors, factor VII, factor VIII and factor IX are also present. Saliva in the mouth also consists of desquamated epithelial cells, microorganisms and their products, few inflammatory cells etc. [2–4].

3.1.2. Formation of saliva

According to Tencate [2], the formation of saliva occurs in two stages. The first stage involves the formation of saliva by the acinar cells. The acinar cells whether serous or mucous cells produce salivary secretion by ribosomal protein synthesis in the rough endoplasmic reticulum

which is followed by the packaging of the proteins by the golgi complex. The secretions are stored as granules and later released into the lumen by the process of exocytosis or by vesicular mechanism. Exocytosis involves fusion of the secretory granules with the membrane allowing release of the contents into the lumen. Vesicular mechanism involves transport of vesicles filled with secretions from golgi complex to plasma membrane. Transcytosis involves passage of substances like immunoglobulin A through the acini. Water is taken up by the cells from the bloodstream and the resulting saliva secreted is isotonic. The serous cells produce serous saliva which is thin, watery and is composed of zymogen granules and contains more proteins, while mucous cells produce thick, viscous saliva containing mucopolysaccharides and mucin. Parotid gland and von Ebner's gland is purely serous gland, while sublingual, glossopalatine and palatine glands have more of mucous secretions. Submandibular gland and other minor salivary gland have both serous and mucous acini, resulting in mixed saliva [2–6].

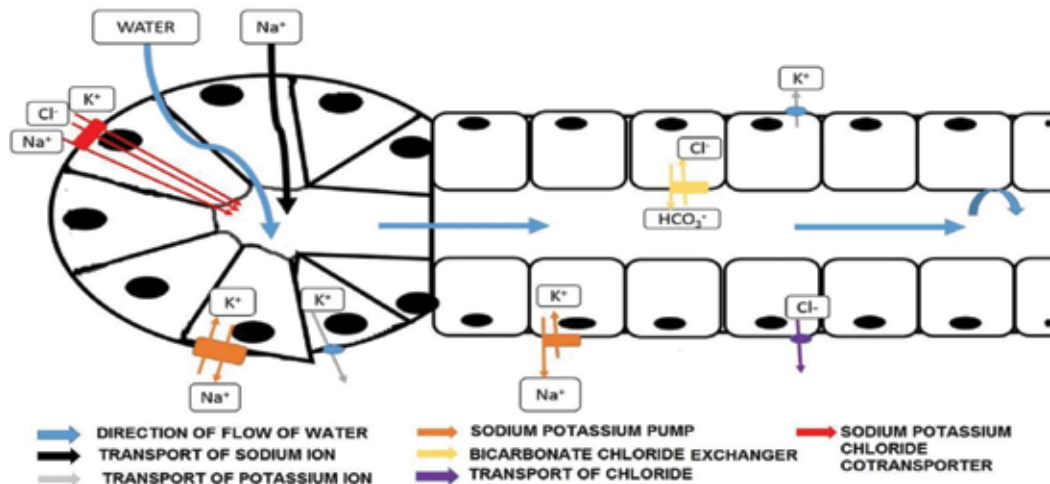


Figure 3. Modification of saliva.

In the second stage the saliva undergoes changes as it passes through the salivary ductal system into the oral cavity. Saliva secreted from the acini is isotonic or slightly hypertonic when it reaches the intercalated ducts. The intercalated duct cells also release lysozymes and lactoferrin. Striated and excretory ducts are impermeable to water. In the striated duct, reabsorption of sodium and chloride occurs more as compared to the secretion of potassium and bicarbonate ions, which makes saliva hypotonic (Figure 3). Striated duct cells also secrete kallikrein and epidermal growth factor. Thus, saliva secreted into the oral cavity is hypotonic as compared to serum [1–6].

3.1.3. Saliva secretion in health and disease

The myoepithelial cells are responsible for the contraction of the acini cells, aiding in the flow and secretion of saliva. In health, the total volume of saliva produced is 750–1000 ml

daily which is contributed by major and minor salivary glands. The resting flow of saliva is 0.2–0.4 ml/min. Salivary flow at rest refers to as unstimulated saliva, whereas salivary flow in response to a stimulus refers to stimulated saliva having a flow rate of 2–5 ml/min. The normal pH of saliva is 6.4–7.4 [1, 2].

A number of factors control the quality and quantity of saliva secreted. The control of salivary gland secretion is mediated by the autonomic nervous system (ANS). All the salivary gland cells receive ANS supply. Control of secretion is also dependent on the perception of taste and smell. The gustatory stimulus is more important than the masticatory stimulus in controlling the salivary secretion. The secretion of saliva occurs by the process of stimulus secretion coupling. This refers to the events involving release of neurotransmitter from vesicles in nerve terminals adjacent to parenchymal cells which stimulate them to discharge secretory granules, water and electrolytes as well as contraction of myoepithelial cells. Norepinephrine activates both alpha and beta adrenergic receptors, while parasympathetic transmitter like acetylcholine activate cholinergic receptors. Alpha adrenergic receptor stimulation results in protein secretion while beta adrenergic or cholinergic stimulation results in low protein secretion and secretion of water and electrolytes. Substance P stimulates alpha adrenergic and cholinergic secretion of saliva. The following flow chart (**Figure 4**) shows the events associated with stimulus secretion coupling which involves the basic process of receptor stimulation which results in increase in the concentration of a secondary messenger, which will further trigger additional events leading to a cellular response [3–6].

Copious watery saliva is secreted in response to parasympathetic stimulation and thicker saliva in response to sympathetic stimulation. Other factors affecting saliva composition are flow rate, circadian rhythm, duration of stimulus, nature of stimulus and diet. During sleep very little saliva is secreted by major salivary glands and majority of the saliva secreted is by the minor salivary glands. Concentration of saliva depends on rate of flow and not on nature of stimulus [2–6].

Historically, it was suggested that parotid salivary gland secretes a hormone called parotin which was considered to have a protein-anabolic function and deficiency resulted in diseases such as chondrodystrophia fetalis, Kaschin-Beck disease, etc. [7].

An increase in the flow of saliva is referred to as sialorrhea (ptyalism), while a decrease in the salivary flow is referred to as xerostomia (dry mouth). Ptyalism is observed after insertion of new orthodontic appliance, in pregnancy, epilepsy, cerebral palsy and Parkinson's disease. Xerostomia is observed in menopause, patients treated by radiation therapy, old age, prolonged use of tranquilizers, amphetamines, antihypertensive and anticonvulsant drugs. A number of systemic conditions affect the functioning of the secretion of salivary glands. Hyperthyroidism, pernicious anemia, vitamin D deficiency, multiple sclerosis and poorly controlled diabetes mellitus affect the salivary glands. Autoimmune diseases like Sjogren's syndrome, Mikulicz's disease affect the salivary gland secretion as the parenchymal elements are affected. Inflammatory, infective and neoplastic diseases also disrupt the activity of salivary gland secretion. Salivary secretion is influenced by hormones. For example antidiuretic hormone facilitates water reabsorption by striated duct, aldosterone causes increased sodium reabsorption by striated duct, testosterone and thyroxine increase salivary secretion [2, 8, 9].

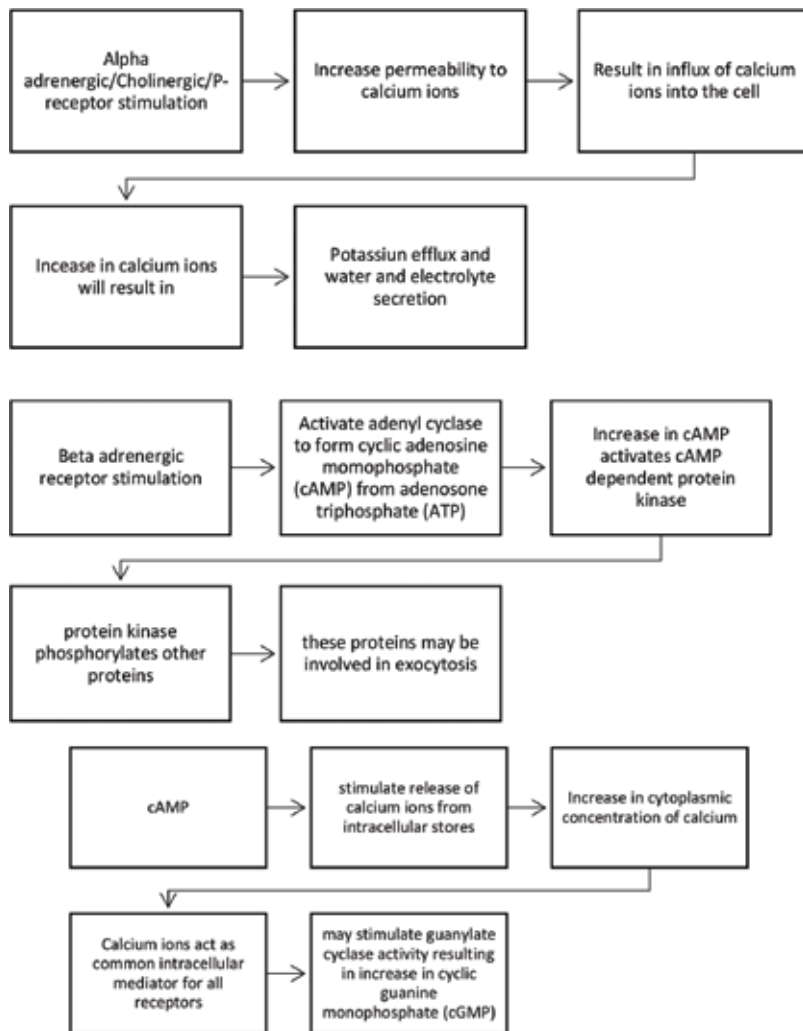


Figure 4. Flowchart depicting sequence of events following neural stimulation.

4. Significance of salivary secretion

The saliva has a number of important functions as mentioned below.

Protection: the saliva contains mucin and glycoproteins which provides it with lubricating properties and moistening the oral cavity, thus preventing friction between the oral structures during physiological functions like mastication. The constant flow of saliva provides clearance of accumulated food debris and microorganisms. Mucins also provide thermal and chemical insulation. Proteins, glycoproteins and mucins form a coating called pellicle formation. Saliva acts as a source of calcium, phosphate, fluoride, statherin and proline rich protein which maintain the integrity of enamel and repair.

Digestion: water and mucin content of saliva aids in bolus formation during the process of mastication. Saliva contains salivary amylase (ptyalin) which helps in digestion of starch and lingual lipase secreted by von Ebner's gland breaks down triglycerides.

Antimicrobial activity: mucins aid in providing a physical barrier to infections by preventing attachment of microorganisms to tooth and tissue surface. Presence of secretory immunoglobulin A provides immune defense. Peroxidase, lysozyme, lactoferrin, histatin, mucins, agglutinin, defensins and cathelicidin also help in providing antimicrobial activity.

Buffering: bicarbonate, phosphate, basic proteins, urea and ammonia help maintain the pH and neutralization of acids.

Tissue repair: salivary glands release growth factors, trefoil proteins into saliva which aid in tissue repair and regeneration.

Taste: saliva acts as a solvent in which molecules from food items can dissolve and reach the taste buds, epidermal growth factor and carbonic anhydrase VI maintains taste buds.

Role of saliva in periodontal pathology: saliva exerts a major influence on plaque initiation, maturation and metabolism. The first step in plaque formation is formation of pellicle followed by plaque formation and maturation [1-6, 8, 9].

Salivary proteins may play a role in plaque mineralization. It is indicated that esterase, pyrophosphatase, acid phosphatase and lysozyme may be involved. Persons with heavy calculus, have higher levels of salivary glycoproteins than non-calculus formers [1-6, 8, 9].

Polymorphonuclear neutrophils (PMNs) reach the oral cavity by migrating through the lining of gingival sulcus. Skougaard and Bay, 1994 believe that orogranulocytic migratory rate correlates with severity of gingival inflammation and is therefore reliable index for assessing gingivitis [8-11].

The saliva acts as an important diagnostic oral fluid owing to its ease and non-invasive mode of collection. A number of components secreted in saliva can be assessed and used to assess diseased states.

A few of the components used as specific biomarkers for detection of periodontal disease include immunoglobulins (Ig) such as IgA, IgM, IgG which interfere in adherence and bacterial metabolism and are present in increased concentration in saliva of chronic and aggressive periodontal patients. Nonspecific markers for aggressive periodontitis include mucins which interfere with the colonization of *Aggregatibacter actinomycetemcomitans* (A. a), lactoferrin which inhibits microbial growth/increased correlation with A. a. Markers for chronic periodontitis include lysozyme which regulates biofilm accumulation and peroxidase which interferes with biofilm accumulation. Nonspecific markers for both chronic and aggressive periodontitis include histatin which neutralizes lipopolysaccharide and enzymes known to affect periodontium and C-reactive proteins which are present in increased concentrations in saliva and serum of patients with periodontitis [8].

Other areas where saliva can be used for diagnosis of diseases and conditions include cystic fibrosis, which is a genetically transmitted disease of children and young adults characterized

by generalized exocrinopathy. In this condition, saliva contains increased calcium levels, elevated levels of sodium and a decrease in flow rate [8, 9].

Sjogren's Syndrome is associated with reduction in lacrimal and salivary secretions. It is characterized by the presence of a lymphocytic infiltrate (predominantly CD4+ T-cells) in the salivary gland parenchyma. A low resting flow rate and abnormally low stimulated flow rate of whole saliva. An antibody p53 can also be detected in the saliva of patients diagnosed with oral squamous cell carcinoma (SCC). Viral diseases like measles, mumps, and rubella can be detected, polymerase chain reaction (PCR)-based identification of virus in saliva is a useful method for the early detection of HSV-1 reactivation in patients with Bell's palsy. Acute hepatitis A (HAV) and hepatitis B (HBV) can be diagnosed based on the presence of Immunoglobulin M antibodies in saliva [8, 9].

Saliva can be used for monitoring of anti-epileptic drugs as a positive correlation between salivary and serum carbamazepine levels has been observed. In another study, salivary levels of phenobarbital and phenytoin demonstrated excellent correlations with serum levels of these medications. Other drugs that can be identified in saliva are amphetamines, barbiturates, benzodiazepines, cocaine, phencyclidine (PCP), and opioids [8–10].

Steroid hormones can be detected in saliva. Salivary cortisol levels were found to be useful in identifying patients with Cushing's syndrome and Addison's disease [12].

Recent focus on the potential role of periodontal disease as a risk factor for cardiovascular and cerebrovascular diseases [13, 14] and the occurrence of pre-term low-birth-weight babies [15] bring new importance to this aspect of salivary analysis [8–15].

Salivary markers as potential diagnostic tests for periodontal disease include proteins of host origin (i.e., enzymes, immunoglobulins), phenotypic markers, host cells, hormones (cortisol), bacteria and bacterial products, ions and volatile compounds [8, 9, 11].

Salivary levels of MMP-8 and IL-1 β appear to serve as biomarkers of alveolar bone loss and hence periodontitis [8, 9, 11].

National Institute of Dental and Craniofacial Research, has highlighted the use of saliva for translational and clinical application by use of salivary proteome and the salivary transcriptome for early detection, disease progression and therapeutic monitoring [8, 9].

Gene therapy has been developed to deliver growth hormone in deficiency states by salivary gland expression of growth hormone [16].

5. Conclusion

The secretions of salivary gland form an integral part of maintaining the physiology of the oral cavity. Saliva is the most important and essential secretion of the salivary glands. Saliva itself has varied functions in the oral cavity and provides additional insight and details of the systemic status of the individual as well. With the advancement in the field of proteomics,

transcriptomics and genomics, better and easier methods of detecting diseases by analyzing saliva are being discovered. For this reason the study of salivary glands and its secretion becomes needful.

Conflict of interest

There is no conflict of interest.

Author details

Anahita Punj

Address all correspondence to: anahitapunj@gmail.com

Army College of Dental Sciences, Secunderabad, Telangana, India

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Xerostomia: An Update of Causes and Treatments

Alejandro Escobar and Juan P. Aitken-Saavedra

Additional information is available at the end of the chapter

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Abstract

Xerostomia or dry mouth sensation is considered a complex condition that affects several stomatological functions that drives to the detriment of the quality of life of individuals who suffer from it. Often, xerostomia is accompanied by a decrease in salivary flow or hyposalivation, and this condition leads to oral health problems such as dental caries, candidiasis, and mucosal complications. Currently, the diagnosis and therapeutic methods for this condition are varied and it is difficult to achieve favorable results in all cases, since the etiology seems to be multifactorial where both local factors and systemic conditions would participate. This chapter presents, in a concise shape, the relevant data about etiology of xerostomia, such as age, autoimmune diseases, systemic diseases, infectious diseases, neuropathic complications, psychogenic factors and therapeutically consumption of drugs among others, and the current available treatments.

Keywords: xerostomia, etiology, diagnosis, clinical manifestation, treatments

1. Introduction

Xerostomia or dry mouth sensation is considered a complex condition that affects several stomatological functions and drives to the detriment of the quality of life of individuals who suffer from it. Often, xerostomia is accompanied by a decrease in salivary flow or hyposalivation with consequences such as oral lesions, alterations of taste, feeling of thick saliva, chewing problems, dental caries, dental demineralization, periodontal disease, salivary gland infection, cervical caries, fungal infections, and others [1]. Currently, the diagnosis and therapeutic methods for this condition are varied and it is difficult to achieve favorable results in all cases, since the etiology seems to be multifactorial where both local and systemic factors would participate [2–5]. Although xerostomia may occur frequently in the general population, clear and defined tools for diagnosis and treatment are still needed. Today, patients suffering from xerostomia visit numerous health professionals to solve this complex condition

that limits many functions of day-to-day life, and often does not find response or effective treatment. Regarding the complexity of xerostomia and its importance in dental practice, this chapter reviews the relevant data about etiology, diagnosis, consequences, and the current available treatments to this condition.

2. Definition and evaluation

Xerostomia (dry mouth, oral dryness, and mouth dryness) is the dryness of oral cavity and can be caused by lower salivary flow or the complete lack of saliva [6]. Based on the etiology, the xerostomia can be classified as true xerostomia (xerostomia vera, primaria), caused by the malfunction of the salivary glands and pseudo xerostomia or symptomatic xerostomia (xerostomia spuria, symptomatica), which is described as the subjective sensation of oral dryness, despite normal secretory function of the salivary glands [7]. The xerostomia, as a symptom, is more common in older populations, but its causes are not related to aging. It has been shown it is related to some specific diseases, drugs, or therapies associated [8]. The prevalence of xerostomia varies from 13 to 28% in older populations, and increases up to 60% in patients living in long-term care facilities [9–11].

Xerostomia, although not considered a disease, may imply the presence of changes directly related to the salivary glands or be the result of systemic diseases [12]. In order for a suitable treatment to be instituted in a timely manner, it is important to carry out a thorough evaluation of the patient with the dry mouth condition, determining, if possible, the cause of the xerostomia. The patients with xerostomia, who are present with salivary gland hypofunction, are at risk of many oral complications; the persistence over time of low rates of salivary secretion causes changes in the oral environment and affects the hard and soft tissues of the mouth. Xerostomia can also be a consequence of systemic disease and its recognition is a valuable aid in the treatment [13]. It is a potentially debilitating condition that can affect up to 1 in 5 oncology patients, with higher prevalence in women and the elderly. There is evidence that the use of multiple medications may increase the risk of xerostomia [13]. This symptom represents a strong impact on the quality of life of the people affected. Over 87.6% of people with xerostomia were worried if they had to spend the rest of their lives with the dry mouth sensation [14]. The dry mouth (xerostomia) sensation has a higher incidence on individuals over 60 years old (12–40%), up to three times higher than on younger adults. It does not seem to be directly related to the normal aging process, but to some chronic diseases or treatments [15, 16]. It is estimated that about 20–30% of the 20-year-old population has xerostomia and the cause may be the increased use of antidepressants, since xerostomia is associated with depression and anxiety. In United States, up to 40% of the 20-year-old population may have xerostomia. The high consumption of antidepressants and other medications, as well as alcoholic beverages and tobacco may explain the increase in people with this condition [17].

Although xerostomia, as a symptom, entails many problems for patients who suffer from it, especially in relation to their quality of life, the decrease in the amount of saliva due to its multiple properties is what brings more consequences at the oral level. Saliva is composed of 99% water and electrolytes. The rest of the composition is organic and includes immunoglobulins,

digestive enzymes such as amylase and lipase, and antibacterial and antifungal enzymes, as well as mucins [14]. Ninety-three percent of its volume is secreted by the major salivary glands and the remaining 7% by the minor glands. Saliva production is controlled by the autonomous nervous system, mainly by parasympathetic nerve signals [18]. Saliva is very important for the preservation of general and oral homeostasis. It has a participation in digestives functions, cleaning, sense of taste, oral mucosa hydration, and defense of teeth trough pH control and its remineralizing potential. In addition, it has antimicrobial properties and controls the composition of oral microbiota by its antibacterial, antiviral, and antifungal capacities [14]. A summary of the Saliva components and functions can be seen in **Table 1**.

Several short and long-term conditions can interrupt salivary secretion, leading to xerostomia. Xerostomia can thus result from three basic causes:

- (1) Factors affecting the salivary center: psychological problems (stress and anxiety), Parkinson’s disease, Alzheimer’s disease (changes in the ability to perceive oral sensations), menopause, and others;
- (2) Factors that alter nerve stimulation of saliva: encephalitis, brain tumors, smoking and dehydration (resulting from the deficiency of water intake, vomiting, diarrhea and polyuria), as well as the use of some drugs, including antihistamines, opioids, antidepressants, antiepileptics, anxiolytics, anticholinergics, antimuscarinics, and others;
- (3) Alterations in salivary gland function as a consequence of obstruction, infection (sialadenitis), glandular tumors, calculi (sialolithiasis), autoimmune diseases (Sjögren’s syndrome-SS-, rheumatoid arthritis, uncontrolled diabetes mellitus and systemic lupus erythematosus), and chemotherapy or radiotherapy performed as cancer therapy for the head and neck area. The extent of injury induced by radiotherapy depends on the volume of irradiated glands and the total dose and technique used [15, 19–23].

Functions	Components
Digestion	Amylase, lipase, ribonucleases, proteases, water, mucins
Phonation	Water, mucin
Taste	Water, gustin
Lubrication	Mucin, proline-rich glycoproteins, water
Antimicrobial action	Lysozyme, lactoferrin, lactoperoxides, mucins, cystins, histatins, immunoglobulins, proline-rich glycoproteins, IgA
Maintaining mucosa integrity	Mucins, electrolytes, water
Cleansing	Water
Buffer capacity and remineralization	Bicarbonate, phosphate, calcium, staterin, proline-rich anionic proteins, fluoride
Preparing food for swallowing	Water, mucins Digestion Amylase, lipase, ribonucleases, proteases, water, mucins

Table 1. Saliva components and functions.

3. Diagnosis of xerostomia

The objective of the diagnosis is to provide treatment as early as possible, thus minimizing side effects in patients suffering from xerostomia. In order to establish a diagnosis of xerostomia, a clinical history is essential to identify the possible etiological factors [24]. It is necessary to investigate its causes. Thus, three orders of factors need to be known: the occurrence of systemic diseases, medication, and the history of radiation therapy. Questions are asked to the patient, trying to find out if he feels the dry mouth sensation, whether he needs to wet his mouth, if he can eat a wafer without drinking water, if the tongue chews the food and clings to the teeth, and the daily water intake daily among other issues [24, 25]. The qualitative clinical diagnosis of xerostomia is made through the observation of clinical signs such as palpation of the salivary glands, observation of the oral mucosa and its hydration, cracked lips, saliva under the tongue, appearance and texture of saliva, the identification of caries, candidiasis and burning sensation, and others [26].

Several methods have been developed to evaluate the level of dryness of the mouth, the discomfort being the most used: sialography, sialochemistry, sialometry and scintigraphy, salivary gland biopsy, ultrasound, magnetic resonance, and computed tomography [19]. Sialography is a technique of imaging that involves the injection of a retrograde form of radiopaque material into the salivary duct system in order to define the anatomy of the glands. This test is very important to demonstrate the presence of nodules or sialectasis, but it has its disadvantages, such as: the difficulty of the technique, since it is invasive and the patient can react acutely or chronically with the contrast material. The biopsy of the major or minor salivary glands allows the detection of inflammatory infiltrations, acinar destruction and dilation of salivary channels with thick mucus, and sometimes fibrosis [27]. Ultrasound, magnetic resonance, and computed tomography are tests that may also contribute to the diagnosis of diseases of the salivary glands.

To establish the condition of the symptom or to evaluate a possible salivary glandular dysfunction, the most used mechanisms are the questionnaire of xerostomia developed by Fox et al. [11, 28] and the determination of salivary flow rate. Sialometry and scintigraphy (an imaging diagnostic method of nuclear medicine that allows the study of the physiology of the various organs) are complementary tests that must be performed in order to evaluate the involvement of the salivary glands in patients with xerostomia. Sialometry is a relatively common procedure in normal clinical practice and include determination of stimulated salivary flow rate (s-SFR), unstimulated salivary flow rate (u-SFR), palatal secretion (PAL), and parotid secretion (PAR). These measurements are the simplest methods of evaluating the salivary glandular function. It is essential to measure the salivary flow, that is, the amount of saliva produced per unit of time. Very low unstimulated and stimulated salivary flow rates or hyposalivation are defined as <0.1 and <0.7 mL/min, respectively [7]. At rest, secretion ranges from 0.25 to 0.35 mL/min and is mostly produced by the submandibular and sublingual glands [29]. Under stimulation, the parotids account for 50% of salivary volume [30]. Determining the stimulated and unstimulated salivary flow is a procedure to measure the amount of saliva it produces a person at a given time. Generally, the stimulated salivary flow is measured for 5 min and unstimulated salivary flow is measured for 15 min [31]. This

kind of measuring has the advantage of being easily implemented, low-cost, and could be available to most of the population at risk [32]. The diagnosis of salivary gland dysfunction is based on data derived from the symptoms reported by the patient, clinical examination leading to verification of the clinical signs and determination of stimulated salivary flow [33]. A severe decrease in salivary flow may lead to a poor health-related quality of life, as well as a risk condition for the development of oral pathologies such as periodontal disease, caries, and candidiasis [29, 34, 35].

4. Causes of xerostomia

The most severe conditions with effect on the salivary flow are SS and radiotherapy in the head and neck area, with the prevalence of xerostomia in almost 100% in these cases. These conditions are characterized by a progressive loss of secretory cells, and thus a progressive decline in saliva production [36, 37]. Less severe conditions may be dehydration, smoking, and inflammation or infection of the salivary glands [12]. In older people, the most common cause of xerostomia is the use of medications because the vast majority of the elderly are being treated with at least one drug that causes salivary hypofunction [32]. A summary of the main causes of xerostomia can be seen in **Table 2**.

4.1. Aging

The reduced salivary flow is commonly seen in the aging populations. This can be attributed to either age-related localized degeneration of salivary glands or systemic diseases [38, 39]. As the patient ages, the organs atrophy and often result in a decrease in salivary production. It was described that in older people the loss was about 30% of acinar cells, with substitution of secretory components by fibrous and adipose tissue [40]. Besides, there are changes in salivary levels of potassium, sodium, IgA, proline-rich protein, lactoferrin, and lysozyme in elderly [28, 40]. A reduction in salivary flow of older people was identified, even in those not using systemic drugs, suggesting a relation between salivary dysfunction and aging [41]. Smith determined that a stimulated salivary flow in healthy adults older than age 70 is lower than in adults under 50 [42].

Systemic diseases	Sjogren's syndrome, diabetes mellitus, Parkinson's disease, encephalitis, brain tumors, Plummer Vinson disease, hypertension, HIV infection, systemic rheumatic diseases, sarcoidosis, Alzheimer's disease, cystic fibrosis, aplasia, chronic tuberculosis, primary biliary cirrhosis, hemolytic anemia, malignant lymphoma, systemic lupus erythematosus, scleroderma, dermatomyositis, pernicious anemia, hypothyroidism, amyloidosis
Other causes of xerostomia (no drugs or systemic diseases)	Radiotherapy and chemotherapy, infections, inflammation, tumors and sialolithiasis in salivary glands, salivary gland excision, vitamin A deficiency, menopause, stress, anxiety, dehydration, neurological disorders, senility, oral sensory dysfunction, iron deficiency, folic acid deficiency, uremia, polyuria, diarrhea, mouth breathing, bone marrow transplantation, endocrine disorders, pancreatic insufficiency

Table 2. Systemic diseases and other causes of xerostomia.

4.2. Drugs

The most common cause of xerostomia is the use of some systemic medications [43]. Several drugs are able to induce hyposalivation and xerostomia, but they rarely cause irreversible damage to the salivary glands. Over 400 medicines, many of them in common use, induce salivary gland hypofunction [44]. Some examples are: anxiolytics, anticonvulsants, antidepressants, antiemetics, antihistamines, antiparkinsonian, antipsychotics, bronchodilators, decongestants, diuretics, muscle relaxants, analgesics, sedatives and anti-hypertensives, and others (**Table 3**) [29]. The exact mechanisms whereby some drugs determine xerostomia or hyposalivation are still unknown. Salivary dysfunction associated to drugs may occur through anticholinergic, cytotoxic action, sympathomimetic, or by damaged ion transport pathways in the acinar cells [39, 45, 46]. Patients who consume a higher number of daily medications have been associated with complaints of xerostomia [47, 48]. The therapeutic and controlled doses of medications do not damage the salivary gland structure. For that reason, drug-induced xerostomia is reversible. The discontinued use of these drugs can restore salivary flow [49].

4.3. Systemic conditions

Xerostomia or hyposalivation may be caused by local factors, including salivary gland disease (sialadenitis) or salivary gland destruction associated with head and neck irradiation for the

Medicine group	Examples
Anxiolytics	Lorazepam, diazepam
Anorectic	Fenfluramine
Anticonvulsants	Gabapentin
Antidepressants—tricyclic	Amitriptyline, imipramine
Antidepressants—SSRI	Sertraline, fluoxetine
Antiemetics	Meclizine
Antihistaminics	Loratadine
Antiparkinsonian	Biperidene, selegiline
Antipsychotics	Clozapine, chlorpromazine
Bronchodilators	Ipratropium, albuterol
Decongestants	Pseudoephedrine
Diuretics	Spirolactone, furosemide
Muscle relaxants	Baclofen
Narcotic analgesics	Meperidine, morphine
Sedatives	Flurazepam
Antihypertensive	Prazosin hydrochloride
Antiarthritic	Piroxicam

Table 3. Medicines and drugs with side effects on salivary secretion.

treatment of cancer [11, 50]. The effects of radiation are dose, time, and field dependent. When the damage of salivary glands by radiation happens is severe [39] permanent gland damage can be expected if the radiation exposure exceeds 50 Gy [50, 51]. Other systemic conditions that also affect the salivary flow are autoimmune diseases (SS, rheumatoid arthritis, AIDS, systemic lupus erythematosus, and scleroderma), neurological disorders (Parkinson's), psychogenic illness such as depression and hormonal disorders (thyroid dysfunction and diabetes mellitus) [9]. Regarding diabetes, we will refer more deeply about it since it is the most frequent metabolic disease in the world and the trend demonstrates that it continues to grow. Both diabetes mellitus (DM) type 1, as type 2 have been associated with xerostomia. In diabetic subjects were shown that salivary flow was significantly lower than in nondiabetic subjects [49]. The causes of low salivary flow may be due to direct injury in the gland parenchyma, changes in the microcirculation to the salivary glands, glycemic control disorders, and dehydration. Some studies consider that this decrease in salivary flow in diabetic subjects is related to an increased diuresis or polyuria, involving a decrease in extracellular fluid and consequently in saliva production [10]. Others explain this as a consequence of dehydration from glycosuria that would be more evident in cases of metabolic decompensation [52]. Regarding neurological diseases, studies have demonstrated that the salivary flow is lower in Parkinson's disease patients. This phenomenon could contribute to dysphagia, which affects up to 75% of patients with this disease [53]. Autonomic dysfunction could explain the decrease in salivary flow in subjects with this disease and the drugs used to their treatment could increase the problem [54]. One of the diseases most associated with a xerostomia is SS, a condition that involves dry mouth and dry eyes and that may be accompanied by rheumatoid arthritis or a related connective tissue disease. The oral manifestations observed in this disease are attributed to the involvement of the salivary glands, which leads to a decrease in salivary secretion [31, 39]. Patients with depression can have hyposalivation medication-induced. However, xerostomia may be of psychological origin. A study observed that subjects with a subjective sensation of dry mouth were significantly more depressive than non-depressive subjects [55]. Another study indicates the possibility of depression as an underlying factor of the sensation of dry mouth [56].

5. Consequences of xerostomia

Patients with xerostomia may have oral and dental consequences. Xerostomia can seriously impact quality of life and may alter speech, eating, and swallowing [13]. The most common complaints of patients with xerostomia include oral discomfort, difficulty speaking, dysphagia, dysgeusia (decreased taste), feeling of thick saliva, and generally, chewing issues, dental caries, dental demineralization, periodontal disease, salivary gland infection (sialadenitis), oral microflora alterations, burning sensation, mucosal inflammation, sore throats, hoarseness, ulcerations, halitosis, mucosal dehydration, reduced lubrication, painful tongue (glossodynia), enlarged parotid gland, oral mucosal fracture, inflammation and fissures of the lips (cheilitis). The reduction of rates of elimination of substances can affect the palate and be associated with changes in the mouth microbiota. The reduction of rates of elimination of substances can affect the palate and be associated with changes in the mouth microbiota. From the mouth, alterations of taste and intolerance to acidic or spicy foods, dry foodstuffs like biscuits can be very uncomfortable for them, and oral cavity examination may exhibit signs such as fissures on the tongue and

lips, angular cheilitis, and dry mucosa. Also, caries, candidiasis, halitosis, or loss of appetite and weight could be observed [25, 57, 58]. This collection of clinical parameters has been indicated as simply estimated for recognizing most patients with xerostomia [38, 47].

The side effects associated with xerostomia are microbial colonization and proliferation in the oral cavity, dental or decreased demineralization, accumulation of stones in the teeth, dehydration of the mucosa, reduction of rates of elimination of substances from the mouth and lubrication of the oral mucosa reduced [13]. When the production of saliva decreases, the buffering capacity of the saliva is reduced, and thus the environment of the oral cavity is vulnerable to acidification, which in addition to determining changes in the normal flora (ecological imbalance) has contributed to the increase in the number of some microorganisms such as *Candida albicans* (a salivary flow less than 0.1 mL/min may cause an increase in the incidence of this fungus) and *Streptococcus mutans* (Gram-negative bacteria). A higher proportion of these microorganisms results in greater acidification of the oral cavity environment, and thus contribute to the enamel demineralization and caries progression. There is a study related to it in which subjects with low salivary flow rate also had significantly more dental caries compared to those with a higher saliva flow rate [58]. In addition, high caries prevalence has been reported to be associated with significantly poorer quality of life compared to low caries prevalence [13].

The infection of the oral mucosa with *C. albicans* affects the lubrication of oral tissues, favoring an increase in the risk of caries and severity of periodontal disease. Candidiasis can also cause burning sensation, glossodynia, glossitis, and angular cheilitis (in areas where the lips are dry or cracked). Patients with prostheses may have reduced retention of the prostheses, pain, and ulcers [59]. The prevalence of oral *Candida* in the normal population has been estimated to range from 23 to 68% and 68 to 100% among SS patients. Studies have attributed the higher prevalence of oral *Candida* carriage in this disease to xerostomia [60].

6. Treatments

Treatment design to alleviate dry-mouth symptoms should be personalized to the individual patient, based on available treatment. The treatments of xerostomia can be classified into the following categories: (1) patient education, (2) prevention, (3) symptomatic treatment, (4) systemic and topical salivary stimulants, and (5) regenerative and gene therapies.

6.1. Patient education

Patients should receive detailed information about the potential causes of dry mouth and the potential sequelae of impaired salivary secretion, such as dental caries, candidiasis, and mucosal complications. Therefore, patients should be encouraged to have preventive oral health care such as dental hygiene habits and regular dental visits [61]. Another palliative action to minimizing symptoms and preventing oral complications is water intake, drinking water frequently, and remaining hydrated is an important treatment for symptoms of dry mouth [1].

6.2. Preventive therapies

Pharmacological interventions for the prevention of radiation-induced salivary gland dysfunction have been studied. The use of chemical radioprotectors represents an obvious strategy to improve the therapeutic index in radiotherapy. However, the vast majority of these are either too weak in terms of radioprotection, too toxic, or without any apparent mechanisms to ensure selective normal tissue protection [62]. The sulfhydryl compound amifostine (WR-2721; 2-[(3-aminopropyl) amino] ethylphosphorothioic acid), is an oxygen scavenger that may protect salivary glands from free-radical damage during radiation therapy without attenuation of the anti-tumor effects in many experiments performed [63]. Amifostine has been approved for prevention of xerostomia, in head and neck squamous cell carcinoma patients undergoing radiotherapy [64]. A recent systematic review that included randomized controlled trials suggested that the drug amifostine prevents the feeling of dry mouth in people receiving radiotherapy to head and neck (with or without chemotherapy) in the short- (end of radiotherapy) to medium-term (3 months after radiotherapy) [65]. However, amifostine has adverse effects such as nausea, vomiting, hypotension, transient, hypocalcemia, and allergic reactions [66]. Then, the benefits of amifostine should be weighed against its high cost and side effects. Another cytoprotective compound described in literature is the bioactive factor Keratinocyte growth factor-1 (KGF-1, also known as FGF-7) [67]. In a phase II Study, recombinant KGF (Palifermin) appeared to reduce mucositis, dysphagia, and xerostomia during hyperfractionated radiotherapy but not standard radiation therapy [68].

Current preventative therapies also include surgical salivary glands relocation outside the radiation field [69]. Jha et al. described a surgical transfer of a submandibular salivary gland to the submental space in order to prevent radiation-induced xerostomia in patients with neoplasias of the pharynx and larynx [70].

6.3. Symptomatic treatment

Saliva substitutes can provide some relief since provide higher viscosity and protection to the oral mucosa [39]. An ideal saliva substitute must simulate natural human saliva, providing long lasting and intense hydration of the oral mucosa, be inexpensive, edible, easy-to-swallow but retainable in the mouth and should allow a minimal number of applications [71]. Saliva substitutes are available in various formulations, e.g., lozenges, sprays, mouth rinses, gels, oils, chewing gums, or toothpastes. Most available in the market contain carboxymethylcellulose (CMC), mucins, xanthan gum, hydroxyethylcellulose, linseed oil, or polyethylene oxide [72]. Subjective impressions of patients suffering from severe xerostomia showed that artificial saliva containing mucins and xanthan gum are better in their rheological and moisturizing properties than those with CMC [73], because mucin-based substitutes had viscosities that were more similar to natural saliva. Recently, it was reported that a polysaccharide-based oral rinse was effective in symptom control in patients with xerostomia and may lead to an increase in saliva production [74]. Other studies include the use of natural products, in this line, a recent double-blinded, placebo-controlled clinical trial, evaluated the efficacy of topical lycopene-enriched virgin olive oil. It showed an improvement of oral quality of life and reduction of xerostomia symptoms [75]. Also, gelatinous substitutes of saliva showed a significant reduction of the

dryness-related complaints in patients suffering from severe xerostomia [76]. A randomized, double-blind, crossover study in patients affected by medication-induced xerostomia showed that two commercial mouthwash plus gel (GUM® Hydral versus Biotène® Oralbalance) achieve a significant improvement in oral health and xerostomia-related quality of life [77]. Recently, a novel edible saliva substitute, oral moisturizing jelly (OMJ), showed a higher grade of satisfaction than a commercially available saliva gel [78]. In addition to the persistent feeling of dry mouth, people who suffer from xerostomia are very susceptible to bacterial, fungal, and other transmittable mouth infections. It is important that products also include human saliva's enzymes (lactoperoxidase, lysozyme, and lactoferrin). Other important feature is to obtain a continuous oral lubrication. In this context, advances in hydrogel technologies and development of buccal mucoadhesive polymers, allows the continuous release of substances that maintain oral hydration and also offer dental-care benefits for its use in treatment of xerostomia [79]. Other strategy involves the use of modified prosthetic structure designed to retain saliva or substitutes in patients who usually wear a dental prosthesis [4, 80].

6.4. Systemic and topical salivary stimulants

Pilocarpine and cevimeline are two systemic US Food and Drug Administration-approved systemic sialogogues for treatment of dry mouth; both can increase secretions and diminish xerostomic complaints in patients, although they must have functional salivary gland cells. Pilocarpine is a cholinergic parasympathomimetic agent that stimulates muscarinic cholinergic receptors on the surfaces of exocrine glands [81] and has been indicated for the treatment of xerostomia [2, 82]. The usual oral dosage for pilocarpine is 5–10 mg three times per day. The initial recommended dose is 5 mg three times per day oral route (OR), which can be increased up to 30 mg/day depending on response and tolerance. The onset of action is 30 min, and the duration of action is approximately 2–3 h. Common side effects include gastrointestinal upset, sweating, tachycardia, bradycardia, increased pulmonary secretions, increased smooth muscle tone, and blurred vision. Contraindications include gall bladder disease, angle closure glaucoma, and renal colic [39, 83]. Cevimeline is a salivary gland stimulant with a stronger affinity for M3 muscarinic receptors [84]. Since it has no effect on M2 receptors, it shows fewer adverse effects when compared to pilocarpine, and besides, it has a long lasting action. The recommended dose is 30 mg three times a day OR, and the most common associated side effect is dyspepsia. Bethanecol is another drug whose action mechanism is on M3 receptors. It has been used to decrease unwanted effects caused by antidepressant and antipsychotic drugs [85]. The dose indicated is four times a day in doses from 10 to 50 mg OR. Adverse effects, despite being infrequent, include nausea and diarrhea. Other drugs that have been put forward include drug with mucolytic properties such as bromhexine improved salivary secretion in patients with SS [86, 87]. Nizatidine, an H2 receptor antagonist alone or in combination with cisapride, showed a significant increase in salivary secretions of dry mouth patients [88, 89]. In addition, other drugs, such as neostigmine, distigmine, yohimbine, nicotinic, and malic acid have also been attributed positive effects in the treatment of xerostomia [3]. Medicinal herbs, such as jaborandi, betel nut, Iceland Moss and Longo Vital, also can stimulate salivary secretion [4].

In the case of tissue autoimmune-related xerostomia, immunologic agents have been used. Interferon alpha (IFN- α), a protein with antiviral and immunomodulating traits, was an

effective treatment for xerostomia linked to SS, improved salivary output and decreased complaints of xerostomia without causing significant adverse medical events [7, 90]. Rituximab (anti-CD20 monoclonal antibody) and infliximab (anti-tumoral necrosis factor—TNF—monoclonal antibody) improved subjective and objective symptoms related to primary SS [91].

Topical salivary stimulants includes sugar-free chewing gum and jellybeans, they can increase salivary secretion by mechanical stimulation and improve the sensation of dry mouth. These products usually contain fluoride, chlorhexidine, calcium phosphate, and xylitol releasers [92, 93], which inhibits the growth of cariogenic bacteria and reduces the incidence of caries [94]. Direct stimulation with electrostimulating device mounted on an intra-oral removable appliance has been used in patients with salivary dysfunction with good results and no significant side-effects [95, 96]. Moreover, non-invasive electrical stimulation systems such as transcutaneous electrical nerve stimulation (TENS) was highly effective in stimulating whole salivary flow in patients with xerostomia and hyposalivation caused by DM and postmenopausal condition [97, 98]. Acupuncture as a method of xerostomia treatment is also cited, a recent randomized and controlled pilot trial of acupuncture showed that acupuncture has beneficial effects on SS symptoms [99]. Other pilot study showed a preliminary evidence that auricular acupressure therapy may be effective in reducing xerostomia intensity in maintenance hemodialysis patients [100].

6.5. Glandular regeneration and gene therapy

Stem cell replacement therapy may be a good option to treat radiation-induced hyposalivation. Stem cell therapy attempts the repair of damaged salivary glands at the cellular level. In this regard, bone marrow stem cells, adipose tissue-derived stromal cells, dental pulp cells have been tested as a form of treatment for hyposalivation after radiotherapy [39]. Interestingly, human salivary stem/progenitor cells (hSSPCs) (derived from parotid and submandibular glands) can be cultured using the salisphere technique and can be introduced to a damaged salivary gland tissue to replace dead or damaged cells. In this context, Pringle et al. showed the presence of SSPCs in cultured human saliphères [101]. These cells were capable of self-renewal and differentiation, which when transplanted into irradiated recipients and restored glandular function. Considering that an ultimate goal is to develop fully functioning bioengineered organs to replace lost or damaged. It was recently reported that a population of SSPCs can be reliably isolated and expanded in sufficient number, suitable for use in a unique 3D hydrogel model of a human implantable salivary gland [102]. However, independent and collaborative work in stem cells research and tissue engineering is still necessary to have fully functional human salivary glands.

Gene therapy involves injecting a vector with genetic information into a tissue to result in some beneficial change. Originally, gene transfer was considered for use in treating congenital genetic disorders, but the basic principles have now been applied virtually to every organ, for acquired as well as inherited disorders. Regarding salivary glands, Baum et al., in phase I/II study, showed an increased saliva flow rate from the targeted parotid gland, as well as a reduction in symptoms related to the radiation-induced xerostomia in subjects treated with the transferring of cDNA for human aquaporin-1 (hAQP1) through an adenoviral (Ad5) vector (AdhAQP1) [103]. Additionally, others genes (Gli1, human keratinocyte growth factor, and Tousled like kinase 1B) have been targeted and have shown promise in preventing salivary

hypofunction in a preclinical mouse [104, 105]. On the other hand, the use of small-interfering RNA (siRNA)-based gene silencing has provided protection of salivary gland from radiation-induced apoptosis at preclinical level [106].

7. Conclusion

Patients with xerostomia are often a challenge regarding diagnosis and treatment, because although xerostomia is not considered a disease, it has a potential devastating effect on the oral cavity. Since dentists are generally challenged with this problem, it is important to have an appropriate comprehension of diverse causes of xerostomia to develop a systematic approach that includes collaboration with physicians to facilitate interdisciplinary patient care, which involves its systemic conditions and medication. Furthermore, a comprehensive management of xerostomia is also necessary and it should incorporate patient education, lifestyle modifications, and adequate pharmacological and non-pharmacological therapies to improve the patient's quality of life. Since most of the successful therapies are depending on the parenchymal gland affection, it is essential to know new therapeutic approaches to fully recover *in vivo* the gland's function or to develop new bioengineered salivary tissues.

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Author details

Alejandro Escobar^{1*} and Juan P. Aitken-Saavedra^{2,3}

*Address all correspondence to: janodvm@gmail.com

1 Institute for Research in Dental Sciences, Faculty of Dentistry, University of Chile, Santiago, Chile

2 Department of Oral Pathology and Medicine, Faculty of Dentistry, University of Chile, Santiago, Chile

3 Post Graduate Program in Dentistry, Federal University of Pelotas, Pelotas, Brazil

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Sialorrhea: A Guide to Etiology, Assessment, and Management

Işıl Adadan Güvenç

Additional information is available at the end of the chapter

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Abstract

Sialorrhea, also known as hypersalivation or ptyalism, is excessive salivation associated with neurological disorders or localized anatomical abnormalities in the oral cavity. Pathologic sialorrhea may develop due to hypersalivation, together with various neurologic disorders including cerebral palsy, Parkinson's disease, and amyotrophic lateral sclerosis, or as an adverse effect of medications. Sialorrhea results in numerous problematic physical and psychosocial complications and has a significant negative impact on quality of life for both the patient and caregiver. The management of sialorrhea is best accomplished with a multidisciplinary team approach. Treatment options range from conservative measures such as observation, positioning, behavioral therapies, and pharmacological therapy to more aggressive methods such as botulinum toxin injections or surgery. The physiology, etiology, assessment, and treatment of sialorrhea are outlined in this review.

Keywords: sialorrhea, hypersalivation, drooling, ptyalism, etiology, assessment, management

1. Introduction

Sialorrhea, also known as hypersalivation or ptyalism, is excessive salivation associated with neurological disorders or localized anatomical abnormalities in the oral cavity. Sialorrhea can be classified as anterior and posterior; both can occur separately or simultaneously. Posterior sialorrhea is the flowing of saliva from the tongue to the pharynx. Anterior sialorrhea results in salivary incontinence or involuntary spillage of saliva over the lower lip, known as drooling. The underlying etiology is the excessive production of saliva or inability to retain saliva within the mouth due to reduced neuromuscular control of the tongue, oral tissues, and

impairment in the swallowing mechanism, all of which are necessary to move saliva from the oral cavity to the oropharynx and beyond [1]. Drooling is common in normally developed babies but subsides between the ages 15 and 36 months with the establishment of salivary continence. Sialorrhea after 4 years of age is generally considered pathologic. Pathologic sialorrhea may develop due to hypersalivation, together with numerous neurologic disorders including cerebral palsy (CP), Parkinson's disease (PD), and amyotrophic lateral sclerosis (ALS), or as an adverse effect of medications. In children, the most common cause of sialorrhea is CP, which persists in 10–38% of these patients. In adults, the most common cause of sialorrhea is PD with a rate of 70–80% [2].

Whatever the cause is, drooling is bothersome, resulting in physical and psychosocial complications. Physical complications include maceration of the skin around the mouth with secondary infection, bad odor, dehydration, speech disturbance, and interference with feeding. People with drooling are also at increased risk for aspiration of saliva, food, or fluids into lungs, particularly when the normal reflex mechanisms such as gagging or coughing are impaired. The psychosocial complications include isolation, barriers to education (such as an inability to share books or computer keyboards), increased dependency and level of care, damage to electronic devices, decreased self-esteem, and difficult social interaction [3]. Sialorrhea may have a significant negative impact on quality of life for both the patient and caregiver [1].

2. Physiology of salivation

The major salivary glands include parotid, submandibular, and sublingual glands; the largest being the parotid gland. These glands secrete saliva, which has a major role in lubrication, digestion, immunity, and maintenance of homeostasis in the human body [2]. The salivary secretion of parotid, submandibular, and sublingual glands is controlled mainly by parasympathetic nervous system, although sympathetic innervation has a minor influence. The parasympathetic fibers originate in the pons and medulla, and they synapse in the otic and submandibular ganglia. Postganglionic fibers originating from the otic ganglion regulate secretory functions of the parotid gland, while the postganglionic fibers from the submandibular ganglion regulate secretory function of the submandibular and sublingual glands. Sympathetic innervation of these glands results in contraction of muscle fibers around the salivary ducts, which enhances the flow of saliva [3].

Salivary secretion is regulated through a reflex arch that produces several actions. The afferent branch contains chemoreceptors in taste buds and mechanoreceptors in the periodontal ligament. Afferent innervations of cranial nerves V, VII, IX, and X carry impulses to salivary nuclei in the medulla oblongata. Efferent impulses are mainly parasympathetic as described above; they come from the chorda tympani nerve (a branch of the cranial nerve VII) and travel to the submandibular, sublingual, and other minor glands via lingual nerve (a branch of the cranial nerve V_3). Efferent fibers to the parotid gland are supplied by lesser petrosal nerve (a branch of cranial nerve IX), which travel through the fibers of auriculotemporal nerve (a branch of the cranial nerve V_3) and reach the gland [2].

The major salivary glands provide 90% of the nearly 1.5 L of saliva that is produced every day. If salivary secretion is not stimulated, that is in basal state, 70% of total salivary secretion comes from the submandibular and sublingual glands. When stimulated, salivary secretion increases by five times, with the parotid glands delivering the greater amount of the saliva [3]. An example of an exogenous source causing stimulation is chewing [2]. There are two main types of saliva produced by the three major salivary glands; serous saliva is produced mainly by parotid glands by stimulation, which is thin and watery, while viscous saliva is produced by sublingual and submandibular glands throughout the day, which is thicker [4]. Both forms of these secretions can be problematic. Serous saliva results in watery saliva, consistently spilling from the side of the mouth, and viscous saliva may be mucoid and sticky, which makes it harder to clear and causes a sensation of choking, associated with panic. It is important to distinguish between thin, runny saliva and thick, mucous secretions because the treatment options differ. The stimulation of cholinergic receptors produces thin, serous secretions, whereas that of beta-adrenergic receptors produces thick protein and mucus-rich secretions. Therefore, in the case of watery saliva secretions, anticholinergics can be preferred, whereas for thick mucus secretions, the addition of beta blockers may be beneficial [5].

3. Etiology of sialorrhea

Sialorrhea associated with neurologic illnesses is generally caused by impaired swallowing due to impaired neuromuscular function. Efficient coordination of various structures, namely the oral cavity, pharynx, larynx, and esophagus, is required for the neuromuscular activity of swallowing. The coordination of these structures forms three phases; the oral phase is under voluntary control, followed by the pharyngeal and esophageal phases, which are involuntary. Spontaneous swallowing is essential for the control of drooling. In children with neurologic disorders, drooling is similarly the result of inefficient tongue and/or bulbar control, thus impaired swallowing, rather than hypersalivation [2].

In children, mental retardation and CP are the most common causes of sialorrhea. Roughly, one in three children with CP is reported to have some degree of sialorrhea. Sialorrhea in CP is caused by oral motor dysfunction, dysphagia, and intraoral sensitivity disorder. Though underestimated, sialorrhea has significant clinical and social consequences concerning the overall health of these children, including dysphagia, respiratory health, and socioemotional aspects of both the children and their caregivers [4]. In adults, PD is the most common etiology. Swallowing impairment, mostly in the oropharyngeal phase, is the major contributor to the pathophysiology of sialorrhea in PD patients, while an increase in the speed of salivary excretion might be a minor contributor. No increase in salivary production was demonstrated in scintigraphic studies in PD patients with sialorrhea, while the speed of salivary excretion of parotid glands in PD patients was significantly higher than normal controls with Tc-99 m scintigraphy [6]. Similarly, in ALS, sialorrhea is not caused by increased production of saliva, but by the inability to swallow secretions because of tongue spasticity, weakness of face, mouth and pharyngeal muscles, and loss of oropharyngeal coordination and function [5]. Less common neurologic causes of sialorrhea are pseudobulbar palsy, bulbar palsy, and stroke (**Table 1**).

Systemic causes

- Neuromuscular/sensory dysfunction—cerebral palsy, Parkinson’s disease, mental retardation, motor neuron disease (ALS), pseudobulbar/bulbar palsy, stroke
- Medication side effects—antipsychotics (clozapine), tranquilizers, anticonvulsants, anticholinesterases, lithium
- Toxin exposure—mercury vapor, pesticides, snake poisoning, mushrooms
- Infection—rabies
- Gastric—gastroesophageal reflux

Local causes

- Oral Inflammation—teething
- Infection—dental caries, oral cavity infection, tonsillitis, peritonsillar abscess
- Anatomic—macroglossia, nasal blockage, oral incompetence, dental malocclusion, orthodontic problems, head and neck surgical defects

Physiological causes

- Pregnancy
-

Table 1. Etiology of sialorrhea.

Increased secretion of saliva frequently develops due to inflammation, such as teething, dental caries, and oral cavity infections. Pregnancy is another significant cause of hypersecretion, usually related to hyperemesis gravidarum. It has an abrupt onset in the 2nd and 3rd week of conception with the rise of hormones and usually resolves during 2nd trimester [7]. Other causes of hypersecretion include side effects from medications (i.e., antipsychotics, tranquilizers, anticonvulsants, cholinergic agonists, and lithium), gastroesophageal reflux, toxin exposure (i.e., mercury vapor, poisonous spider bites, mushrooms, insecticides), and rabies [3, 8]. Clozapine, an antipsychotic used in schizophrenia, is a rather common cause of sialorrhea, which manifests in 30–80% of patients taking the drug. Hypersalivation usually develops early in the treatment course and is typically more prominent at night [9].

Anatomic abnormalities are usually not the only cause of sialorrhea; however, most of the time, they exacerbate other causative conditions. Macroglossia (enlarged tongue) and oral incompetence may cause salivary spilling. A constantly open mouth due to nasal blockage or malocclusion and other orthodontic problems may compound oral incompetence; treatment of nasal problems and orthodontic correction can alleviate sialorrhea. Surgical defects occurring after major head and neck surgeries may result in sialorrhea as well [3].

4. Assessment of sialorrhea

Assessment of the severity of sialorrhea and its impact on the quality of life for the patient and the caregivers assist in establishing a prognosis and appropriate management of the problem. History should be taken from the patient and the caregiver to understand the etiology and severity of the situation and its impact on the daily life. Use of medications, language and communication skills, cognition, respiratory health, and presence of gastroesophageal reflux disease should be questioned. In physical examination, oral cavity should be examined for sores on the lip and chin, dental problems, tongue size and movement, and tonsillar

Drooling severity	Points
Dry (never drools)	1
Mild (wet lips only)	2
Moderate (wet lips and chin)	3
Severe (clothing becomes damp)	4
Profuse (clothing, hands, tray, objects become wet)	5
Drooling frequency	
Never drools	1
Occasionally drools	2
Frequently drools	3
Constantly drools	4

Table 2. Drooling frequency and severity scale.

hypertrophy; nasal blockage, malocclusion, and jaw stability should be assessed. A neurological examination should be carried out investigating the level of alertness, swallowing ability, motor skills, and sensory dysfunction of the patient. The nutrition and hydration status, head posture, and emotional state of the patient should also be evaluated [4, 10].

Objective and subjective measures have been developed to quantify sialorrhea. The objective test methods include radioisotope scanning, collection cups strapped to the patient's chin for the measurement of salivary flow, and direct observation of saliva loss such as counting the number of napkins used daily to contain excessive saliva production, measuring the weight of the towels or dental cotton rolls [4, 10]. The importance of objective methods is that they seem to be more sensitive in detecting a reduction in sialorrhea or drooling than purely subjective assessments [11].

A variety of subjective scales for sialorrhea have been described. Subjective scales such as the drooling frequency and severity scale, the drooling rating scale, the drooling impact Scale, and visual analog scales can be given to patients or their caregivers to determine the qualitative and quantitative consequences of the severity and impact of sialorrhea [4]. The drooling frequency and severity scale is an easy comprehensive scale, which rates the severity of drooling on a five-point scale and the frequency of drooling on a four-point scale (**Table 2**) [12]. Subjective scales are useful and appropriate methods to measure changes in sialorrhea, because the impact on families, caregivers, and the patients themselves is of utmost importance when assessing satisfaction with the effectiveness of any treatment [4].

5. Management of sialorrhea

The management of sialorrhea continues to be a challenge in spite of various effective treatment strategies to diminish saliva production. The flow of saliva from the oral cavity to the esophagus depends on numerous factors, such as cognitive and mental abilities, intact swallowing, oral sensibility, lip closure, and ability to keep the head upright [1].

Treatment of sialorrhea is best accomplished with a multidisciplinary team approach. The complete history and physical examination of the patient, the assessment of the impact of drooling on quality of life, and the potential for improvement can be undertaken by the primary care physicians. Speech pathologists and occupational therapists provide education for swallowing mechanics to the patients and support their posture with devices such as the head back wheelchair. Dentists and orthodontists identify and correct dental and oral diseases and malocclusion. Otolaryngologists diagnose and treat causes of aerodigestive obstruction like macroglossia and adenotonsillar hypertrophy that contribute to drooling. Neurologists assess the severity and prognosis of neurologic conditions that result in drooling [3].

The goal of the treatment of drooling is a reduction in excessive salivary flow, while maintaining a moist and healthy oral cavity. Avoidance of xerostomia (dry mouth) is essential. The two main approaches are:

1. Noninvasive modalities including positioning, improving eating and drinking skills, oral facial facilitation, speech therapy, biofeedback, positive and negative reinforcement, oral prosthetic devices, pharmacological therapy, and botulinum toxin
2. Invasive modalities including surgery and radiotherapy

Generally, no single approach is adequately effective, and usually, a combination of therapies is used. Primarily, reversible causes of drooling should be treated. Less invasive and reversible methods are preferred before surgery is undertaken [10]. Behavioral approaches and therapies employed by speech pathologists are rarely curative, while systemic medications and surgical approaches may have severe and long-term adverse effects [1].

For minimal problems, in children younger than 4 years of age or in adults with unstable neurologic function, observation may be the convenient choice. Minimal issues can be handled with a feeding program aimed at improving oral-motor control as well; nevertheless, this can rarely be helpful. Anatomical problems should be identified and treated, and adenotonsillectomy should be undertaken, if necessary. Dental malocclusion and caries should be corrected. Patients should be fitted with appropriate wheelchairs and braces, when required. A number of orthodontic appliances may be used, such as customized plates that fit the palate for improving lip closure or movable beads placed on the upper plate that stimulate tongue movement, thus helping to deflect saliva toward the pharynx [3].

Other conservative therapeutic options include positioning techniques, oral-motor, and speech therapies given by speech therapists, which improve oral awareness and motor control. Biofeedback and automatic cueing techniques may be utilized in patients with mild neurologic dysfunction and drooling. These devices are used to associate a behavior with a cue, such as swallowing or wiping the face with a beep sound. Reinforcement methods, which are suggested behaviors such as encouraging patients for swallowing and wiping their faces and discouraging open mouth, can be used as an adjunct in moderate sialorrhea [3, 10].

Whenever sialorrhea continues to affect the patient's health and quality of life in spite of these conservative measures, pharmacological therapy and other invasive therapies should be considered.

5.1. Pharmacological therapy

Oral therapy for sialorrhea encompasses the use of anticholinergic agents such as glycopyrrolate, benztropine, scopolamine, and tropicamide. Anticholinergic agents work by downregulating acetylcholine and ultimately decreasing saliva secretion through the parasympathetic autonomic nervous system. Glycopyrrolate oral solution is an anticholinergic agent that was the first drug approved in the United States for drooling in children with neurologic conditions and is generally well tolerated. However, anticholinergic agents are poorly tolerated by elderly patients. Glycopyrrolate actually has lower risk of central side effects, owing to its quaternary ammonium structure, that makes it impossible to pass the blood–brain barrier in large amounts. It is effective and safe at 1 mg, 3 times a day [2]. Studies have shown 70–90% response rates, but approximately 30–35% of patients discontinue the drug due to side effects such as excessive dry mouth, urinary retention, decreased sweating, skin flushing, irritability, and behavior changes [10]. Other undesirable adverse effects observed with such treatment include constipation, urinary retention, tiredness, and drowsiness [11]. Anticholinergics are contraindicated in patients with glaucoma, obstructive uropathy, gastrointestinal motility disorders, and myasthenia gravis.

Intraoral tropicamide films provide short-term relief of sialorrhea. One study provided evidence that 1 mg of tropicamide resulted in significant visual analog scale score decrease and reduction in saliva volume in nondemented PD patients [2]. Transdermal scopolamine, applied as a patch behind the ear, was well tolerated in short-term studies, but its use was limited by side effects of urinary retention and blurred vision [3].

A comprehensive systematic review of the use of anticholinergics in children concluded that benztropine, given 3–3.8 mg per day, could be effective. A significant decrease in the mean score for drooling was reported with benzhexol hydrochloride (2×2 up to 2×3 mg daily). There was also some evidence for a marked decrease in drooling with glycopyrrolate [13]. Benztropine was also reported to show a significant reduction in the total salivation scores compared to botulinum toxins A and B in a study of mixed treatment network meta-analysis of randomized controlled trials on pharmacological interventions for treating sialorrhea associated with neurological disorders [14].

Antireflux medication has also been suggested for use in drooling; however, there are no double-blind studies in the literature to offer evidence for this recommendation [2].

5.2. Botulinum toxin

The injection of botulinum toxin (BT) to the major salivary glands has grown in popularity because of its limited invasiveness and demonstrated effectiveness in many patients. It has been shown to improve quality of life of patients effectively with a low profile of side effects. However, it is important to take into account that the duration of the therapeutic effect is limited in time, generally lasting a few months [11].

The effect of BT in sialorrhea was first reported in PD patients [15]. This toxin is a potent neurotoxin that blocks the release of acetylcholine and a number of other neurotransmitters from synaptic vesicles; hence, it shows its effect by blocking cholinergic postganglionic parasympathetic fibers in sialorrhea [2].

Currently, three type A and one type B toxin are approved for use in the US. These are OnabotulinumtoxinA (BOTOX®), AbobotulinumtoxinA (Dysport®), IncobotulinumtoxinA (Xeomin®), and RimabotulinumtoxinB (Neurobloc®/myobloc®) [2]. Both A and B types of BT are reported to be effective in treatment of sialorrhea, and both have a low profile of side effects [1].

BT can be injected in parotid and/or submandibular glands. The dose, concentration, and volume of injectate, number of injections, injection site, rate of injection, gauge of needle, and distance of needle tip from the neuromuscular junction are among the factors that can affect the diffusion and spread of BT, thus its efficacy in sialorrhea. A broad dose range of BT has been reported in various studies, specifically from 10 to 100 U of Botox®, from 20 to 300 U of Dysport® per patient, while usually 2500 U of Neurobloc® per patient is reported to be injected. The effect of BT on salivation lasts for 1.5–6 months [11]. Older age is significantly associated with longer benefit duration [16]. Sometimes the reduction in salivary secretion and improvement in drooling may not be correlated, owing to the variability of the factors that influence the severity of drooling and reduction of saliva secretion [2]. Patients with PD showed a more favorable safety-efficacy ratio than did patients with ALS, due to lower adverse events and longer benefit duration [16].

In a recent meta-analysis, eight randomized placebo-controlled trials involving 181 patients were reviewed. The study reported that BT improved drooling severity in patients with sialorrhea significantly in both adult and pediatric populations. Increased saliva thickness (3.9%), dysphagia (3.3%), xerostomia (3.3%), and pneumonia (2.2%) were reported as common side effects [1]. Adverse effects such as chewing difficulties and recurrent mandibular luxation have been reported [11]. BT therapy is reported to have many advantages over other noninvasive and invasive treatments. It is effective and minimally invasive with few side effects and a low risk of aspiration. However, it is expensive and temporary, and the need for repeated sedation can be troublesome with children [2].

Ultrasound guidance for intraglandular injection is preferred by some of the authors. Blind puncture of the superficial lobe of the parotid gland following anatomical landmarks is easier, because the structure is relatively superficial. Since the submandibular glands are normally nonpalpable, infiltration may be more challenging. Ultrasound easily identifies the glandular structures for infiltration, while avoiding accidental damage to other anatomical structures, the facial nerve in the case of injection into the parotid gland, or the facial vessels in the case of the submandibular gland [11].

Jongerius et al. compared the efficacy of BT injections to transdermal scopolamine. They reported that even though both treatments were successful in significantly lowering drooling parameters, patients treated with BT did not experience any side effects, while 40% of patients taking scopolamine reported severe adverse effects. The wide range of side effects and potential drug–drug interactions encountered with scopolamine and glycopyrrolate suggests that BT may be a safer option compared to systemic anticholinergics [13].

BT can also be used for empirical selection of patients who would, in the future, be good candidates for surgical treatment of the major salivary glands. In this way, patients who respond

well to BT injections can be treated more efficiently with surgery rather than receiving multiple injections. On the other hand, patients who do not respond well to BT may be considered as poor candidates for surgical treatment, because failure rates could be much higher owing to the contribution from minor salivary glands to the etiology of sialorrhea [17].

5.3. Radiation therapy

Radiation therapy to the salivary glands is a useful treatment option in elderly patients who are not candidates for surgery and cannot tolerate medications. Radiation causes xerostomia that lasts months to years. The dose may be changed to produce the desired effect, and it can be repeated if required. The main problem is that radiation can induce malignancies, but this does not happen until 10–15 years after treatment and therefore are less of a concern in patients who are elderly and debilitated [3].

5.4. Surgical treatment

While many patients are successfully treated with conservative methods and medical therapies, a number of patients are not able to tolerate the side effects of medications. BT treatment is reported to show improvement in drooling, but surgery provides a larger and longer lasting effect. Surgeons should consider more aggressive interventions for patients with chronic sialorrhea secondary to neuromuscular dysfunction with the impairment of swallowing. In these cases, sublingual or submandibular gland excision, submandibular duct ligation, parotid duct ligation, submandibular or parotid duct rerouting, or any combination of the above procedures result in higher rates of success, both short term and long term, and they may be cost-effective compared to BT injections requiring multiple visits [18]. Nevertheless, it is important to mention that surgery has a risk of permanent consequences (especially xerostomia), and that it should be preferred only in severe cases who are not responsive to nonsurgical therapies and in whom sialorrhea has great impact on the health and quality of life of the individual and caregivers [5].

Tympanic neurectomy is now regarded as a historical technique used to denervate salivary glands. This technique is performed through the middle ear, where the tympanic plexus and chorda tympani travel before entering the major salivary glands. The procedure is relatively simple and fast, but salivary function returns within 6–18 months, when nerve fibers regenerate [3, 17].

Recently, a novel procedure, transoral endoscopic submandibular ganglion neurectomy, was performed in two cases of BT-resistant drooling. Six months follow-up was successful; however, long-term results are awaiting to be warranted [19].

The most definitive treatment of sialorrhea is to excise the major salivary glands or to ligate or reroute the major salivary ducts. Surgical management can be described by a combination parotid duct ligation or rerouting with either submandibular gland excision or submandibular duct rerouting. Preservation of salivation with decrease in drooling could be accomplished by rerouting of the parotid and submandibular ducts to the posterior oropharynx with the advantage of no external scar. There may be a potential for aspiration after these procedures. Sublingual gland

excision is suggested as well when the submandibular ducts are rerouted to prevent formation of salivary retention cysts. Parotid duct ligation is a simple fast procedure without an external scar, which decreases the stimulated salivary flow. There may be a risk of sialocele development. The most definitive surgical procedure, which includes bilateral parotid duct ligation and submandibular gland excision, is highly successful, with nearly total elimination of sialorrhea, a low incidence of facial weakness, and significant patient and caregiver satisfaction [3].

The meta-analysis of surgical management using a variety of surgical procedures demonstrates significant subjective relief in 81.6% of pediatric patients with sialorrhea following surgery. Bilateral submandibular gland excision and parotid duct rerouting had the highest reported success rate of 87.8%. However, simple bilateral submandibular duct rerouting and bilateral submandibular duct rerouting with bilateral parotid duct ligation had similar levels of subjective success. Four-duct ligation had the lowest success rate at 64.1%. Although this procedure seems simpler and less invasive compared to other surgeries, it can cause significant pain and swelling, since the ligated glands continue to produce saliva for a period before atrophy occurs. Bilateral submandibular duct rerouting is a procedure that is more complex than a simple submandibular duct ligation. The reported success rates with this procedure are consistently good and similar to procedures involving submandibular gland excision. Limited data suggest that fibrosis of the gland occurs due to obstruction of the rerouted duct, ending up as an actual ligation of the duct, but in most of the patients, function is maintained in at least one gland [17].

Author details

Işıl Adadan Güvenç

Address all correspondence to: isiladadan@yahoo.com

Department of Otorhinolaryngology, Head and Neck Surgery, Cigli Regional Education Hospital, Izmir, Turkey

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New Approaches in the Diagnosis of Salivary Gland Diseases

Salivary Diagnostics

Karolina Elżbieta Kaczor-Urbanowicz

Additional information is available at the end of the chapter

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Abstract

Salivary diagnostics plays an important role in the early detection and prevention of many oral and systemic diseases in a fast and noninvasive way. Saliva collection is an easy, repeatable and inexpensive diagnostic source that can be used for both diagnosis and real-time monitoring of various human diseases. In the near future, many developed and validated salivary biomarkers have the potential to reach the clinical practice. Five diagnostic “omics” constituents of saliva include proteomics, transcriptomics, metabolomics, microbiomics and microRNAs. Based on them, the newly emerging technologies of salivary diagnostics are developed that include RNA-sequencing, point-of-care technologies and liquid biopsy. They have potential to enable screening, early detection, prognosis and monitoring of various human diseases. The recent developments broadened the salivary diagnostic approach from the oral cavity to the whole physiological system, thus toward personalized individual medicine applications.

Keywords: saliva, diagnostics, oral cancer, RNA-sequencing, point-of-care, liquid biopsy

1. Introduction

Saliva is an encouraging medium to be used in the early detection, diagnosis and monitoring of oral and systemic diseases, specifically for the purpose of personalized medicine by incorporating point-of care technology platforms in the clinical settings. Though, saliva collection is easy, fast, cheap, safe, does not require specialized equipment and can be performed at home [1]. The normal daily production of saliva varies between 0.5 and 1.5 l. Saliva is an acidic biofluid, derived from the three major salivary glands (parotid, submandibular, sublingual) as well as from minor glands (labial, buccal, lingual, and palatal tissues). It is composed in vast majority of water (99%), while other constituents occur in trace amounts, including proteins and both inorganic (sodium, potassium, calcium, magnesium, chloride, etc.) and organic

constituents (amylases, peroxidase, lipase, mucins, lysozyme, lactoferrins, cystatins, hormones, etc.) [2]. Salivary composition is very diverse (RNA, DNA, proteins, metabolites, and microbiota), and may be utilized for diagnostic purposes. Most importantly, salivary components may vary in their concentrations and levels depending on the individual's health or disease status. Thus, real-time monitoring of salivary data can provide useful translational clinical applications in the detection of various human oral and systemic diseases. The development of the recent technologies based on salivary diagnostics will help to introduce screening programs to enable early detection and monitoring of the disease [3].

Saliva is an important biofluid with lots of various biological functions including lubrication, chewing, swallowing, sensation, digestion and protection of oral mucosa against biological, mechanical, and chemical factors as well as infections [2].

2. Salivaomics

Currently, there are known five major diagnostic toolboxes of saliva "Salivaomics": proteomics (the study of proteins), transcriptomics (the study of RNAs), metabolomics (the study of metabolites), microRNA (the study of microRNAs) and microbiome (the study of microbiota) [1, 2] (**Figure 1**).

2.1. Proteomics

The proteomics is the large-scale screening for proteins, their expression, modifications, and interactions by using high-throughput approaches [4, 5]. Proteins can indicate various physiological and pathological states of the current health condition or specific disease. The recent advancements in proteomics contributed to the development of new non-invasive technologies.

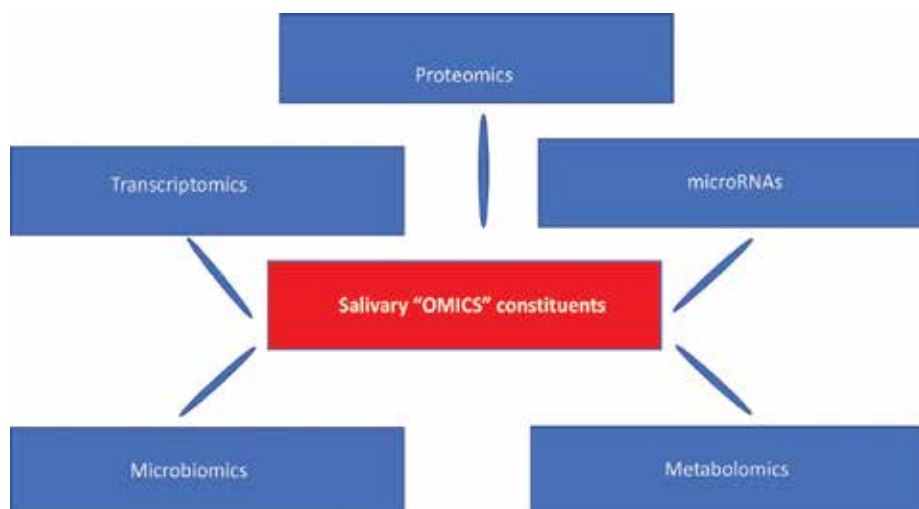


Figure 1. Diagnostic toolboxes of saliva.

The currently accepted gold standard methods for proteomic analyses include: triple depletion of high abundance proteins (removal of albumins, alpha amylase and immunoglobulins), sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS PAGE), two-dimensional gel electrophoresis (2-DE), mass spectrometry (label free qMS), ELISA (enzyme-linked immunosorbent assay) and Western blotting [2, 6]. Other advanced methods include electrospray ionization (ESI), matrix-assisted laser desorption ionization (MALDI), quadrupole/linear ion trap, time-of-flight (TOF), quadrupole TOF (QTOF), Fourier transform ion cyclotron resonance (FT-ICR), or the OrbiTrap, MS/MS, MALDI-MS or targeted HPLC-ESI-MS/MS [7].

Recently, a great focus has been put on identification of salivary protein biomarkers for various human oral and systemic diseases such as: pancreatic cancer, Sjogren's syndrome, oral cancer, lung cancer, orthodontically induced root resorption, etc. [2, 5]. The proteomics delivers an alternative ideal and non-invasive diagnostic tool, more sensitive, and safer for detecting the disease status. In addition, the depletion of high abundance proteins from saliva contributes to significant increase in the detectability of less abundant salivary proteins [5, 8]. There are known three major methods of high-abundance protein removal [9]: enzyme-substrate absorption method used for alpha-amylase affinity removal [8], immunodepletion method and combinatorial peptide ligand library (CPLL) [10].

Proteomic analysis of saliva is commonly used in the diagnostics of oral diseases as well as general health disorders such as oral candidiasis [11], oral squamous cell carcinoma (OSCC) [12], glossodynia [13], head and neck squamous cell cancer [14], Sjogren's syndrome [15], HIV [11], autism [16], fibromyalgia [17], breast cancer [18], lung cancer, melanoma [19] or pancreatic cancer [7].

Various mediators associated with oral cancer are released from cells due to malignant conditions and have been analyzed in saliva samples, like cytokines, chemokines, interferon-gamma (IFN- γ), interleukins (IL-1 β , IL-6 and -8, IL-4 and -10), tumor necrosis factors (TNF- α), transforming growth factor-beta-1 (TGF- β 1), epidermal growth factor (EGF), vascular endothelial growth factor (VEGF), and endothelin [20]. In case of oral squamous cell carcinoma, elevated levels of NF- κ B-dependent cytokines have been observed in saliva [21]. Other potential protein biomarkers include IL-6 and S100A9 [22] or BGH3, MMP9 and PDIA3 [23]. In addition, increased salivary expression levels of MMP1 and MMP3 in OSCC patients can indicate more advanced stage of disease [12], while adenosine deaminase might be indicative of early stage of oral tongue cancer [24].

2.2. Transcriptomics

Transcriptomics (gene expression profiling) is the quantitative study of an organism's transcriptome, all RNA transcripts present in a cell. The information is recorded in the genome and expressed through transcription. These data can be used for capturing marked changes in expression levels of specific genes in the detection of various human diseases [2, 25]. Transcriptomics encompasses a great diversity of RNA species including messenger RNAs (mRNAs), long non-coding RNAs (lncRNAs), and small RNAs such as microRNAs (miRNAs), transfer RNAs (tRNAs), piwi-interacting RNA (piRNA), etc. The mRNAs play an important

role in carrying information for making proteins while noncoding RNAs have different functions [25]. In turn, the microRNAs (miRNAs) are a group of small non-coding RNAs that regulate mRNA through sequence-specific binding to the UTR [26]. The miRNAs are involved in various biological processes.

The common practice in identification of salivary transcriptomic biomarkers is microarray technology. However, this technology is currently replaced by the newer one, RNA-sequencing (RNA-Seq).

2.2.1. RNA sequencing

RNA-sequencing (RNA-Seq) has been the newly developed method for characterizing the full human transcriptome. RNA-sequencing (RNA-Seq) has a wide variety of applications, but no single analysis pipeline can be used in all cases and for all biofluids [27]. Specifically, RNA-Seq of saliva is challenging, including difficult RNA isolation step to enhance yield of salivary exRNA step as well as laborious RNA-Seq small and large library construction stage, inclusion of spike in standards and controls, RNA-sequencing, data storage and data analysis. Therefore, currently available literature regarding salivary RNA-Seq is scarce [28–30].

RNA-sequencing is a rapidly progressing transcriptome profiling that uses deep-sequencing technologies and becoming the major tool in analyzing gene expression [31]. This is a new high-throughput method for both mapping and quantifying transcriptomes. It provides more detailed information about the levels of transcripts and their isoforms than other methods [32]. In general, a total RNA is converted to a library of cDNA fragments with adaptors attached to one or both ends. Afterwards, a sequencing of the molecule is performed, with or without amplification, that provides short sequences from one (single-end sequencing) or both ends (pair-end sequencing). The reads are typically 30–400 bp, depending on the DNA-sequencing technology used. Revealing the transcriptome is crucial for investigating the functional elements of the genome, the molecular constituents of cells and tissues, and also for understanding development and disease [32].

2.2.1.1. Advantages of RNA sequencing

RNA-Seq (RNA sequencing) has clear advantages over existing approaches and is believed to become the best method for analyzing transcriptome in the near future.

Firstly, RNA-Seq can be used for detection of both known (corresponding to existing genomic sequences) and novel transcripts, thus enabling identification of new organisms with unidentified yet genomic sequences. RNA-Seq is used for precise localization of transcription boundaries, to a single-base resolution. Furthermore, this method can be used for examining transcripts of great complexity as it provides useful information about the connectivity of exons as well as sequence variations in the transcribed regions [33–34].

Secondly, RNA-Seq has very low, if any, background signal. This feature differentiates it significantly from microarray platforms as it can be uniquely mapped to the genome regions of interests [35].

RNA-Seq has also significant benefits over array-based technologies for detecting expression quantitative trait loci (eQTLs). Though, it can identify different transcript variants and enable quantification of allele-specific expression within an individual to increase association mapping [27, 36, 37].

Finally, it has a highly accurate and large dynamic range of expression levels with high reproducibility rates while using less RNA sample and at a much lower cost than either tiling arrays or large-scale Sanger EST Sequencing [33, 35].

2.2.1.2. Disadvantages of RNA sequencing

The major problems of RNA-Seq technology are associated with RNA isolation and cDNA library construction as it includes several manipulation stages, which can complicate profiling of transcripts of various lengths [32]. This causes the variability in the measurements, influenced by the technical noise [27].

In addition, longer RNAs have to be fragmented into smaller pieces (200–500 bp) to be compatible with most deep-sequencing technologies using RNA or cDNA fragmentation methods (RNA hydrolysis or nebulization, DNase I treatment, etc.) [33, 35]. These sequencing procedures include a number of steps (RNA fragmentation, cDNA synthesis, adapter ligation, PCR amplification, bar-coding, and lane loading) that might introduce biases into the resulting data [27].

Another key consideration concerning library construction is whether or not to construct strand-specific libraries [33, 38], that provide information about the orientation of transcripts, essential for transcriptome annotation. However, strand-specific libraries are very time-consuming to produce [39, 40].

RNA-Seq has also bioinformatics challenges associated with storage, retrieval and processing large amounts of data. The alignment of long RNA-Seq reads is also complicated due to non-unique mapping to multiple locations in the genome [32].

Also, to detect a rare transcript or variant, more sequencing depth is needed. In general, the larger the genome, the more sequencing depth is required for adequate coverage, which brings greater cost [32].

Although RNA-Seq is still in the early stages of use, it has clear advantages over previously developed transcriptomic methods such as microarray profiling [32]. Specifically, salivary RNA biomarker transcripts of IL8, IL1B, DUSP1, HA3, OAZ1, S100P, and SAT yielded high sensitivity (91%) and specificity (91%) in distinguishing OSCC from the controls [41]. Also, salivary IL6 mRNA and IL-8 may serve as potential biomarkers for diagnosis of OSCC [42, 43].

2.2.2. Difficulties in salivary RNA-sequencing and bioinformatic analysis

RNA-sequencing (RNA Seq) of saliva is challenging compared to other biofluids such as blood or urine. RNA Sequencing of salivary samples is associated with several problems such as inadequate technique of RNA isolation, improper stabilization of RNA or RNA library construction [44–48]. Till now, there were also no established guidelines how to bioinformatically process the salivary RNA-Seq data. The recent paper by Kaczor-Urbanowicz et al. gives the

recommendations for bioinformatics analysis of salivary RNA-Seq data that differs from other biofluids (blood, urine, etc.) as saliva contains the majority of microbial content, while other physiological fluids are considered to be sterile [48]. Thus, it is recommended to use quite stringent and sensitive criteria, while working with salivary RNA-Seq data to avoid erroneously mapped bacterial reads to the human genome, and to prevent problems with their further annotation to human RNA databases. In addition, the specific sequence of alignment steps and the stringency parameters associated with processing of RNA Sequencing data can grossly increase the final data quality [48].

2.3. Micro-RNA-omics

MicroRNAs (miRNAs) are short, single-stranded RNAs that are about 21 nucleotides in length. Their function is to regulate gene expression. Like other types of RNA, miRNAs are transcribed from DNA, but they do not participate in protein translation. They are non-coding RNAs, in which each primary transcript (pri-miRNA) is processed into a pre-miRNA and finally into functional miRNA [49]. Mature miRNAs are involved in various biological processes such as cell growth, differentiation, apoptosis, stress and immune response or glucose secretion [50–52]. Studies on miRNA dysregulation in various human diseases have risen rapidly in recent years, including those in cancer, heart disease as well as type II diabetes mellitus and its complications, such as endothelial and vascular smooth muscle cell dysfunction, cardiomyopathy and nephropathy [53–55]. Most importantly, salivary microRNAs (miRNAs) (miR-9, miR-134 and miR-191) can be used as potential biomarkers for head and neck squamous cell carcinoma [56]. The reduction in salivary expression profiles of miR-125a and miR-200a was observed in OSCC patients compared to healthy people [57]. In turn, miR-31 increases in OSCC patients, specifically in saliva, where it rises even more than in plasma [58].

2.4. Metabolomics

Metabolomics is the study of small molecular metabolites of living tissues, mostly metabolic intermediates such as carbohydrates, lipids, amino acids, nucleic acids, etc. [1]. The major metabolomic technologies include high-performance liquid chromatography-mass spectrometry (HPLC-MS), two-dimensional gas chromatography MS and nuclear magnetic resonance spectroscopy in conjunction with pattern recognition methods [2].

Salivary metabolites are involved in many biological processes as well as pathogenesis of various diseases such as periodontal diseases, renal diseases, hepatocellular carcinoma and colorectal cancers [59] as well as oral cancer [60]. In case of oral leukoplakia, an upregulation of putrescine, 8-hydroxyadenine and 5,6-dihydrouridine in OSCC can be indicative of increased risk for malignant transformation [61].

2.5. Microbiomics

Microbiomics include study of bacteria, archaea, protists, fungi and viruses. Microbial profiling (Human Oral Microbe Identification Microarray) of salivary microbiome in early resectable pancreatic cancer revealed that *Neisseria elongata* and *Streptococcus mitis* were successfully

developed with 96.4% sensitivity and 82.1% specificity [62]. Currently, newer microbiome-based technologies have been developed such as RNA or DNA sequencing [1]. In addition, two microbial biomarkers, Firmicutes (especially *Streptococcus*) and Actinobacteria (especially *Rothia*) were significantly decreased in oral cancer compared to healthy controls [63]. Finally, Furquim et al. reported that patients with Fanconi anemia (FA) are at higher risk of developing OSCC than the general population, especially after the hematopoietic stem cell transplantation [64].

3. Modern technologies in salivary diagnostics

3.1. Salivary liquid biopsy

Recently, a new trend appeared to reveal emerging role of “liquid biopsy” as identification method of biomarkers in various cancers. Liquid biopsy tests are non-invasive biofluid tests (serum, urine, saliva) that detect circulating tumor cells (CTCs) and fragments of tumor DNA shed into the bloodstream by cells undergoing apoptosis or necrosis [3].

The role of liquid biopsy markers including circulating tumor cells, circulating RNAs (miRNA, lncRNAs and mRNAs), cell-free proteins, peptides and exosomes has been currently investigated as non-invasive cancer biomarkers in different biofluids such as blood, urine, saliva and seminal plasma. Liquid biopsies hold great promise for personalized medicine due to the fact that they enable multiple non-invasive global sampling resulting in longitudinal assessment of the primary and metastatic tumors. Molecular profiling of circulating molecules (proteomic, transcriptomic, genomic, metabolomics, microRNAs) contributed to the successful application of several non-invasive multi-marker tests in the clinic [65].

Nowadays, liquid biopsy enables a variety of clinical and investigational applications such as early detection, assessment of molecular heterogeneity of general disease, monitoring of tumor dynamics (in melanoma, breast, ovarian or colon cancers), identification of genetic determinants for targeted therapy, evaluation of early treatment response, monitoring of minimal residual disease or assessment of resistance evolution in real time [66].

The most common technologies of liquid biopsy include detection and quantification of ctDNA (circulating tumor DNA) in blood such as Sanger sequencing, pyrosequencing, next generation sequencing, PCR-based technology, high-performance liquid chromatography (HPLC), mutant-enriched liquid chips, amplification refractory mutation system (ARMS), beads, emulsion, amplification and magnetics (BEAMing), pyrophosphorolysis-activated polymerization (PAP) [2, 66] or electric field-induced release and measurement (EFIRM) [67, 68]. The current gold standard methods for detection of ctDNA targets include droplet digital PCR and next-generation sequencing. However, those technologies require extraction of DNA from large volume of biofluid samples. EFIRM can be successfully used for continuous monitoring during treatment. The results are very promising [3].

Circulating tumor DNA (ctDNA) is considered to be stably found in biofluids encapsulated in extracellular vesicles (EVs) and released by cells into the circulation. If the links between distal

cancers and the oral cavity will appear to be scientifically proven, it will open a new avenue of clinical utility to effectively, and non-invasively diagnose cancers through saliva. The ctDNA mutant fragments were observed in plasma [69, 70] and saliva samples [71] of head and neck cancer patients.

3.2. Point-of-care technologies

The current knowledge of salivary biomarkers and their role in point-of-care applications highlights the need for development of more advanced technologies. As a consequence, point-of-care diagnostics is definitely approaching reality for salivary research and closely related with its translation into clinical practice [3] as it delivers information of the current status of the disease in a very fast, convenient and non-invasive way. PoCs can be successfully used for early detection and real-time monitoring of the disease [3].

The current PoC technologies are ubiquitous. They comprise microfluidics, micro/nanoelectromechanical systems (MEMS/NEMS), paper-based technology, RNA-sequencing, liquid biopsy, biosensors, fluorescent biosensors, photometric and electrochemical methods, electronic nose and electric field-based methods such as electric field-induced release and measurement (EFIRM) method [3, 68, 72]. Contemporary available PoCs can be delivered in form of small and portable smartphones or “lab-on-chips” [3].

One of such PoC development is the Oral Fluid NanoSensor Test (OFNASET), that is used for multiplex detection of salivary proteomic (thioredoxin and IL-8) and genomic biomarkers (messenger RNA biomarkers, i.e. SAT, ODZ, IL-8, and IL-1b) for oral cancer with 90% sensitivity and 90% specificity for both interleukin 8 (IL-8) and IL-8 protein messenger RNA (mRNA) [67]. In turn, OraRisk human papilloma virus (HPV) test with Reflex (Quest Diagnostics, Los Angeles, CA, USA) can be indicative of HPV infection, high risk factor for development of oral cancer [68]. In addition, electrical controlled magnetic EC Sensor is designed to detect microRNA-200a [73], electrochemical sensor using endonuclease target recycling amplification to capture oral cancer overexpressed 1 (ORAOV1) [74], while wireless mouthguard enzymatic biosensor to detect uric acid [75] or lactic acid [76], potential biomarkers for oral cancer.

4. Conclusions

Salivary diagnostics is a promising field for the implementation of PoC technology. The desire for PoC, the potential of saliva, development of validated panel of salivary biomarkers for specific diseases and development of novel advanced techniques enables the application of saliva for the early detection and diagnosis of several oral and systemic diseases in a non-invasive, easy and fast personalized way. The recent technology advances, including liquid biopsy, EFIRM, biosensors, smartphones, microfluidics, paper-based technology, have the potential to make clinical utilities of saliva a reality in the near future. Saliva is predicted to be a substitute for blood, collected non-invasively for the diagnosis of oral and systemic diseases as well as chairside screening.

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Conflict of interest

The author reports no conflict of interest in relation with the present study.

Notes/thanks/other declarations

Nothing to declare.

Author details

Karolina Elżbieta Kaczor-Urbanowicz

Address all correspondence to: kaczor.karolina@gmail.com

Department of Oral Biology and Medicine, Division of Oral Biology, UCLA Center for Oral/Head and Neck Oncology Research, UCLA Laboratory of Salivary Diagnostics, UCLA Department of Orthodontics, University of California at Los Angeles (UCLA), California, USA

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Proteomics of the Salivary Fluid

Goran Mitulović

Additional information is available at the end of the chapter

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Abstract

Following the sequencing of the human genome, the mapping of the human proteome is the next task to be completed in order to gain knowledge on how proteins are involved in disease genesis, growth, therapy, and healing. As contrary to the genome, which is relatively static, the human proteome is significantly more complex and highly dynamic. Whilst the majority of the research is being focused on analyzing either the proteome of tumor tissues and tumor cells or the proteome of serum and plasma, little attention has been awarded to the analysis of proteomes in saliva or urine. The proteome in saliva can help providing important information on processes involving health issues in dentistry, head and neck cancers, gastric cancers or neurology, to name just a few. However, this is changing and the proteomics research community is increasingly focusing on deciphering the salivary proteome. So far, more than 3000 proteins have been identified in different studies and more is to come with new instrumentation and methods available. Some of the proteomics methods applied for analysis of salivary proteins will be discussed in this chapter.

Keywords: salival, proteomics, diagnostics, biomarkers, chromatography, mass spectrometry

1. Introduction

The raise of proteomics and the continuous development and improvement of analytical instruments such as high-performance liquid chromatography (HPLC) and mass spectrometry (MS) have substantially fuelled the development of instrumental methods for analysis of proteins for both research and clinical questions [1–6]. Proteomics is not only addressing the efficient separation of peptides upon, mostly, tryptic digestion of proteins and their sensitive detection using mass spectrometry. Proteomics is a technology enabling significantly and profoundly better approach to investigating and understanding proteins' function and their posttranslational modifications [7–12]. Applying proteomics methods for analysis of clinical samples is especially important in time of personalized medicine, which tailors individualized treatment of each

patient based on specification of the diseases [3, 13–19]. Furthermore, (clinical) proteomics can be used as a method-of-choice for the screening of biomarkers used for early discovery and early diagnostics. Early diagnostics and early discovery, needless to say, will help decreasing patients' morbidity and mortality by detecting the disease at the stage when it can be effectively treated with less side effects and at significantly lower cost for the society. This approach can be very beneficial especially for diseases affecting large populations such as cardiovascular diseases, diabetes mellitus and other endocrinal diseases, glioblastoma and similar.

Proteomics can also be applied for the point-of-care diagnostic approaches where both medical professionals and patients can get rapid information and bed-side diagnosis. Of course, classic proteomics approaches with protein extraction from tissue or body fluids and overnight protein digestion cannot be applied; however, proteomics can provide information to be used with kits for point-of-care approach [20–24].

It has already been shown that saliva is a highly valid biological fluid that can be used for diagnostic applications [25–39].

Various number of components can be identified in saliva, which provides real-time data on the patient's condition. The substances found in saliva include but are not limited to DNA, RNA, proteins, metabolites, and microbiota from both oral and gastrointestinal origin. Sample collection is simple, cheap, and can be provided by patient at home without expensive equipment or medical personal needed on-site.

This manuscript will provide a short insight into different techniques applied for proteomics analysis of saliva starting with sample collection, protein precipitation and digestion, peptide separation and MS detection, and finally with data analysis.

2. Sample collection

For medical or biological analysis in general, the method how and when a sample is being collected is of utmost importance. Sample collection is one of the pre-analytical steps that need extraordinary caution and that can influence, badly in most cases, the complete process of sample analysis.

Saliva is mainly composed of water; however, there are a number of other substances being present. Mucins, proteins, DNA, RNA, enzymes, sugars, cell debris, and microbiota and their secretome can interfere with test performance. Therefore, an optimized sample collection process is needed, and a researcher must carefully and cautiously prepare the sample collection step, and the patient must be educated and trained if sample is being to be collected at home. Furthermore, steps like sample storage and transport to the laboratory must also be carefully planned and executed especially when longitudinal studies of the same patient or of the patients' groups are being performed, and especially in parts of the world where appropriate infrastructure is not always available [40–45].

Several methods can be applied for collection of saliva: (1) passive droll is the simplest approach but the saliva often has a high concentration of mucins and high viscosity and (2) the Salivette[®] Systems [46–48], the Greiner Bio-One Saliva Kit, and the recently introduced

RNA-Prosal [49]. All three sample collection systems have been used in the field, and publications describing their efficacy are available [50–52].

At the Proteomics Core Facility of the Medical University of Vienna Salivette® is being used for induced saliva sample collection by chewing cotton swabs. As mentioned previously, it is of great importance to carefully plan and perform sample collection. The patient or the donor must refrain from consuming food, alcoholic beverage, and caffeine at least for 2 h before sample collection. Further, the patient shall briefly wash the mouth using water only. Saliva is being collected for 2 min during which the patient chews the cotton swab. This approach yields approximately 2.5 ml of saliva, which is sufficient for performing proteomics analysis. Some patients, however, need additional stimulation for saliva production and paraffin gum can be used in these cases to stimulate saliva flow and gain enough sample volume. In these cases, the use of Greiner Bio-One Kit helps obtaining more saliva than chewing the cotton swab; however, one shall be careful since this kit contains citric acid, which can lead to protein denaturation and protein loss during the sample collection.

As soon as the sample has been collected it shall be supplemented with enzyme inhibitors in order to suppress enzyme activity and protein degradation. A total protease inhibitor cocktail such as Roche’s “Complete Protease Inhibitor Cocktail®” is being added to the sample following centrifugation and removal of cellular debris and prior to storage at –80°C.

It is of extreme importance to secure reproducible sample collection procedures and properly train the patients in cases of self-sampling to avoid sample contamination and alteration. Furthermore, conditions for proper sample transportation and handling until it is being processed must also be carefully considered and applied.

Table 1 shows the steps applied for sample collection and the preparative work.

Step	Device	Temperature	Precautions
1. Sample collection	Salivette® (used in the Proteomics Core Facility at the Medical University in Vienna)	Human body	No food, alcoholic or caffeine beverages until 2 h before sample collection!
2. Centrifugation	Centrifuge	4°C	
3. Protein precipitation	Modified Wessel-Fluegge as described or Aceton	According to the protocol described	Sample should always be prepared on ice!
4. Depletion of high abundant proteins	Antihuman serum albumin and Anti-IgG columns and anti-amylase column	Ambient to 40°C	Pay attention to columns’ loadability!
5. Enzymatic digest	Offline digest overnight	37°C	
6. (Multidimensional) HPLC separation and MS detection of tryptic peptides	Nano HPLC and mass spectrometer	Various combinations	Column capacity, compatibility of selected separation dimensions
7. Bioinformatics-Database search	Various platforms are available	Not relevant	Carefully select parameters, avoid very stringent but also very lax conditions
8. Verification and Validation			

Table 1. General description of sample preparation for proteomics analysis of salivary samples.

3. Sample preparation: depletion of abundant proteins and enzymatic digest

The enzymatic digest of salivary proteins does not differ from digestion of other proteins. Usually, trypsin is being used for proteolytic cleavage of proteins due to its relatively high specificity, availability, and ease of use. Furthermore, tryptic peptides are ideally suited for reversed phase HPLC separation and positive ionization using electrospray MS (ESI). Tryptic peptides are also ideal for using multidimensional separation approaches such as strong cation exchange (SCX), hydrophilic interaction liquid chromatography (HILIC), or electrostatic repulsion interaction chromatography (ERLIC) since they bear positive charges on N-terminus or Lys-residues.

Beside trypsin, other proteases can also be used alone or in combination with trypsin. The aim is either to achieve specific cleavage of proteins for special questions or to achieve smaller peptides and enhance their ionization, detection, and ultimately better sequence coverage for identified proteins. Mostly, proteins such as Lys-C and GluC are applied for pre-digestion before trypsin addition.

Protein digest begins with protein precipitation from saliva. A number of methods have been developed for protein precipitation such as: (1) alcoholic; (2) salting out; (3) applying strong acids (trichloroacetic acid); (4) using acetone; (5) using acetonitrile, etc. Protein precipitation shall help for removing DNA and RNA and their fragments from the sample and for removing lipids.

The Proteomics Core Facility of the Medical University in Vienna applies a modified Wessel-Fluegge [53] method for protein precipitation [54]. **Figure 1** shows the sample collection steps prior to protein reduction, alkylation, and addition of the protease.

In addition to trypsin, the use of proteases, e.g. LysC, will help generating smaller peptides prior to separation and detection [55, 56].



Figure 1. Sample collection steps prior to protein reduction, alkylation, and addition of the protease as performed at the Proteomics Core Facility of the Medical University of Vienna (<https://www.sarstedt.com/en/products/diagnostic/salivasputum/product/51.1534.500/>).

Unlike serum or plasma, saliva does not contain large range and amounts of high abundant proteins such as serum albumin or hemoglobin that can affect sensitivity and selectivity of detection. However, amylase and serum albumin are still proteins with the highest abundance in saliva and can also affect the detection of other low abundant species and should be removed from the sample [57, 58]. In human saliva, alpha-amylase makes about 60% of the abundance of all proteins present, and its removal will help identifying proteins of lower abundance such as cytokines, which can be used as putative biomarkers for different processes. Deutsch et al. [57] have shown a simple yet very effective method for removing alpha-amylase and gaining a deeper insight into saliva's proteome. The use of a simple potato starch resulted in sixfold reduction of the amount of alpha-amylase in the sample. Albumin removal can be facilitated by using a number of columns developed and based on immunoaffinity reactions [59].

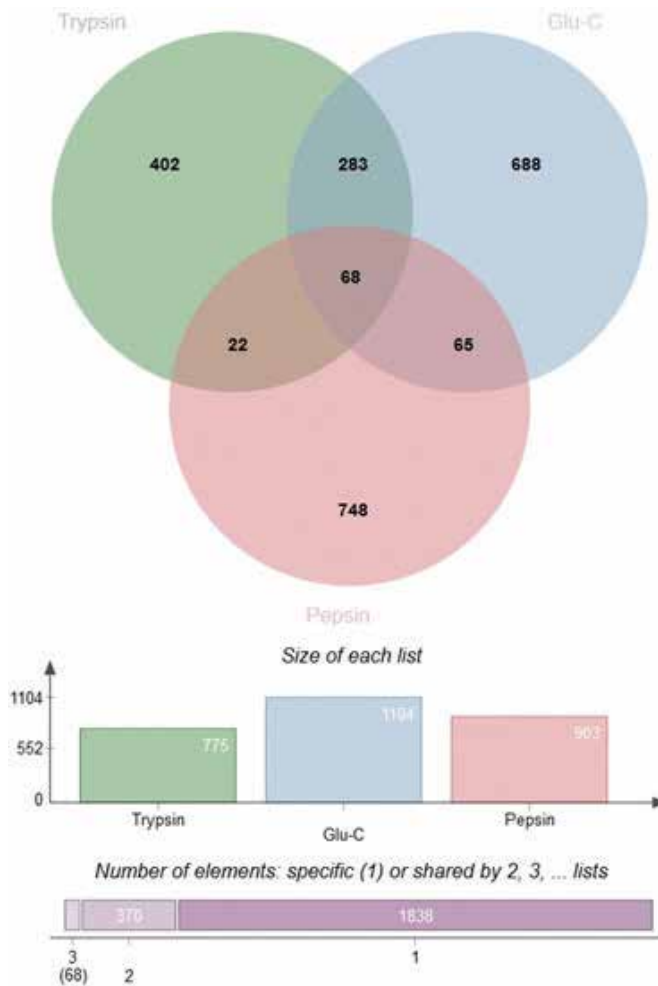


Figure 2. Comparison of the number of identified proteins from human saliva by applying different proteases. Courtesy of Zofia Świąteczak (Master Thesis).

Enzymatic digest of salivary proteins does not differ from enzymatic digests used for other biological samples. Based on previous experience and results, trypsin is the most widely used protease for protein digest resulting with peptides suited for subsequent separation using cation exchange columns or anion exchange columns, reversed phase separation, and mass spectrometric detection. A comparison of results obtained using different digestion methods for salivary samples obtained from healthy donors is shown in **Figure 2**.

As shown, the highest number of proteins was identified upon applying a combined digestion approach and using GluC and trypsin. However, the choice of proteases used also depends on analytical problem to be addressed as, e.g. for detection of glycosylated proteins, which might require additional proteases to be applied.

4. Chromatographic separation of digested proteins

Upon tryptic digestion of proteins, resulting peptides are being separated on a chromatographic column prior to mass spectrometric detection and subsequent bioinformatics analysis. Separation of peptides is being performed either using one-dimensional approach or the multidimensional separation by combining two or more separation technologies prior to MS detection and analysis.

For the one-dimensional chromatographic separation approach, peptides are being injected onto the reversed phase nano HPLC column where they are separated according to their hydrophobic interaction with the stationary phase [60]. **Figure 2** shows an exemplary base peak chromatogram (BPC) for one-dimensional analysis separation of salivary peptides. Usually, long separation gradients are selected for one-dimensional separation in order to provide the best possible conditions for peptide separations and large number of identifications.

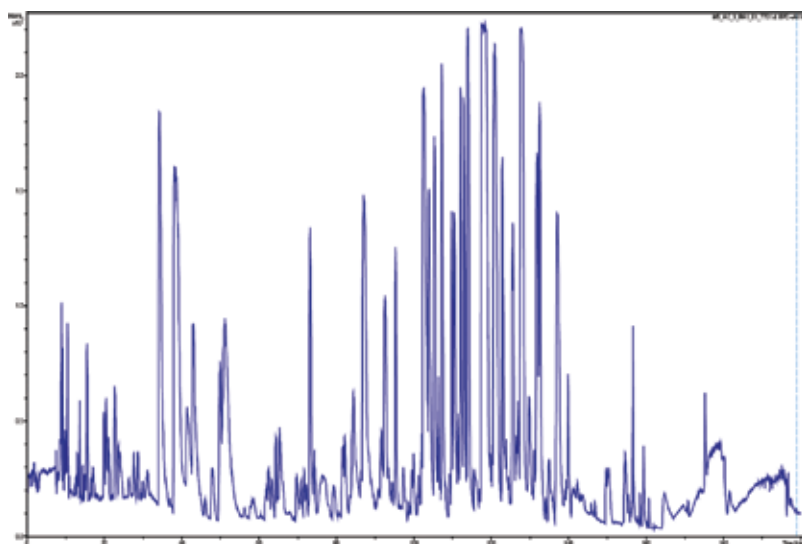


Figure 3. Base Peak Chromatogram (BPC) of GluC-tryptically generated peptides, which were separated on a reversed phase nano HPLC column.

The separation shown in **Figure 3** was performed using a 180-min gradient and a total analysis runtime of 210 min. Thus, the total amount of available time must be considered when performing this kind of analysis.

The use of multidimensional separation methods will increase the number of peptides detected and the number of identified proteins, and, in addition, protein's sequence coverage will be

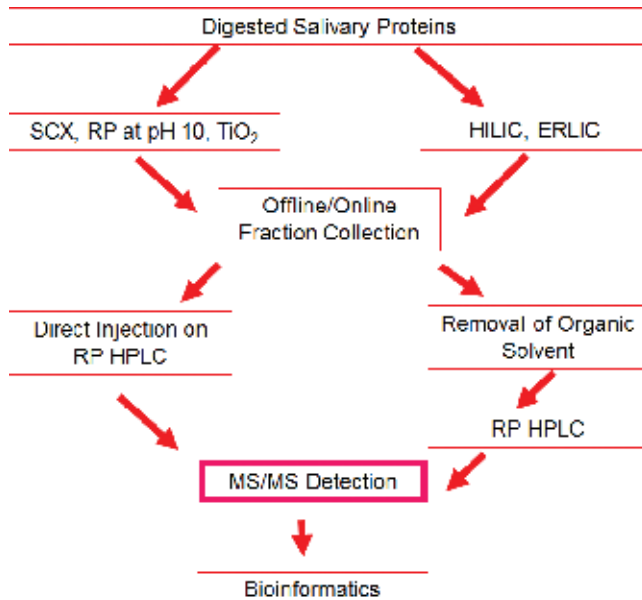


Figure 4. A number of combinations of different techniques can be used for separation of peptides and proteins in a proteomics approach.

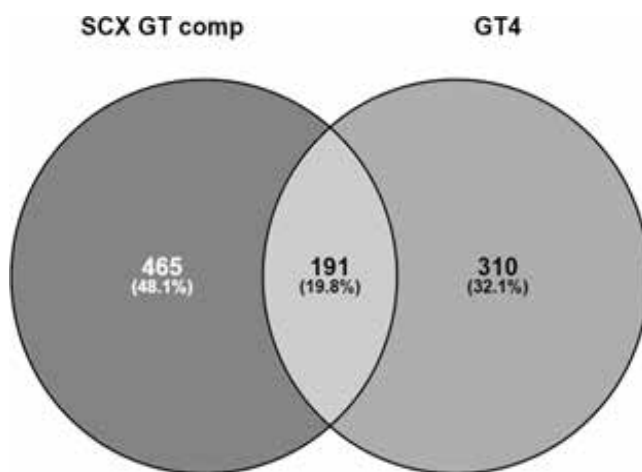


Figure 5. Two-dimensional separation approach for salivary peptides enables identification of higher number of proteins. Data courtesy of Zofia Świątczak (Master Thesis).

improved for identified proteins. Improved sequence coverage for identified proteins is one of the major challenges for proteomics analysis with high importance since it enhances chances for detection of posttranslational modifications (PTM) such as phosphorylation, glycosylation, methylation, etc., which are important as drug targets.

Different types of separation approaches can be used for the two-dimensional approach: strong cation exchange columns, weak anion exchange columns, reversed phase columns at high pH, and other combinations are possible [61–63]. A schematic of possible combinations of chromatographic approaches is shown in **Figure 4**.

Generally, the use of multidimensional separation will result in increased number of identifications, and **Figure 5** shows the comparison of the number of detected proteins upon applying the two-dimensional chromatographic separation with strong cation exchange column used for the first separation dimension.

5. Mass spectrometric detection and bioinformatics analysis

Upon separation, peptides are being detected using mass spectrometry and analyzed by comparing experimental data and databases of in-silico digested proteins. Several MS approaches have been applied for detection of salivary peptides: Electrospray Time-of-Flight (ESI-ToF), MALDI-Time-of-Flight (MALDI-ToF), ESI-Orbitrap analysis, ESI-Quadrupole ToF, etc.

Depending on MS type and selected instrumental method, posttranslational modifications of proteins can also be identified and thoroughly analyzed thus enabled a deeper insight into the proteome. The majority of top-down analysis, i.e. analysis of undigested proteins is performed using MALDI mass spectrometers, and the majority of analysis for digested proteins (peptides) is performed using electrospray (ESI) ionization and ToF and Orbitrap mass analyzer.

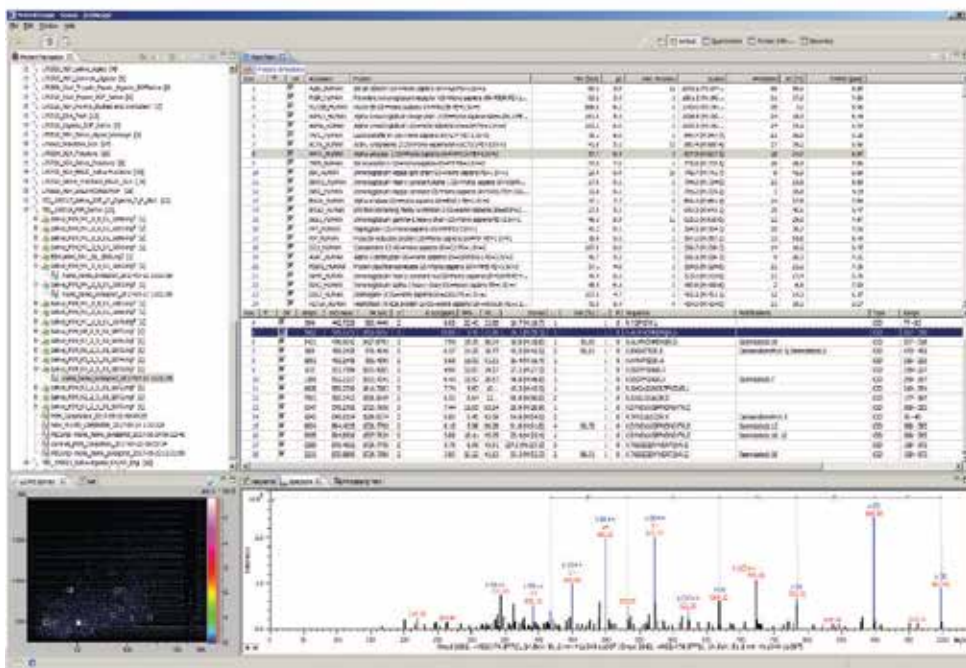
The analysis of obtained raw data is performed by searching protein databases such as SwissProt, Uniprot, NR (by NCBI), and user-generated databases. A number of commercially available software packages such as Mascot (Matrix Science, London), ProteinScape (Bruker, Germany), Proteome Discoverer (Thermo Scientific, Bremen, Germany), and of free available software such as The Global Proteome Machine (www.thegpm.org), MaxQuant (<http://www.coxdocs.org/doku.php?id=maxquant:start>), PeptideShaker (<http://www.uib.no/en/rg/probe/65218/peptideshaker>), Skyline (<https://skyline.ms/project/home/begin.view?>), OpenMS (<https://www.openms.de/>), and other packages. The choice of the software to be used strongly depends not only on personal preferences but also on data to be analyzed and the information needed to be extracted.

Figure 6 shows a screenshot of two software packages preferably used at the Proteomics Core Facility at the Medical University of Vienna, Peptide Shaker and ProteinScape.

In addition to database search and protein identification, the analysis of the pathways where proteins are being over- or underexpressed and the analysis of interactions with other proteins have been performed using free software such as DAVID[®] (<https://david.ncicrf.gov/>), STRING (<https://string-db.org/>), Reactome (<http://reactome.org/>) or commercially available MetaCore[®] (<http://lsresearch.thomsonreuters.com/>) or similar.



Peptide Shaker



ProteinScape

Figure 6. Screenshots showing analysis of a salivary sample by applying two distinct software packages. Note that identified proteins have been listed based on their scores, which can be calculated using different algorithms.

Term	Count	Genes	Count	%	Adjusted	ES
Salivary secretion	37	37	1.2	6.3E-9	8.4E-6	
Glycolysis / Gluconeogenesis	27	27	0.9	4.5E-6	6.0E-3	
Carbon metabolism	35	35	1.2	9.6E-5	1.3E-1	
ABC transporters	18	18	0.6	2.7E-4	2.9E-1	
Glutamateric synapse	34	34	1.1	2.7E-4	3.6E-1	
Biosynthesis of antibiotics	54	54	1.8	3.3E-4	4.3E-1	
Amoebiasis	32	32	1.1	3.4E-4	4.5E-1	
Oocyte meiosis	32	32	1.1	5.0E-4	7.7E-1	
Calcium signaling pathway	45	45	1.5	1.5E-3	1.9E0	
Platelet activation	34	34	1.1	3.2E-3	4.2E0	
Pathogenic Escherichia coli infection	17	17	0.6	4.2E-3	5.4E0	
Biosynthesis of amino acids	22	22	0.7	4.4E-3	5.7E0	
Regulation of actin cytoskeleton	49	49	1.6	4.9E-3	6.4E0	
Vascular smooth muscle contraction	31	31	1.0	5.3E-3	6.9E0	
Chemokine signaling pathway	44	44	1.5	5.6E-3	7.2E0	
Circadian entrainment	26	26	0.9	5.9E-3	7.5E0	
Endocytosis	57	57	1.9	7.2E-3	9.2E0	
Renin secretion	19	19	0.6	8.8E-3	1.1E1	
Excretory secretion	25	25	0.8	8.9E-3	1.1E1	
Pentose phosphate pathway	11	11	0.4	1.1E-2	1.3E1	
Arrhythmogenic right ventricular cardiomyopathy (ARVC)	20	20	0.7	1.3E-2	1.6E1	
miR signaling pathway	24	24	0.8	1.3E-2	1.6E1	
Tight junction	33	33	1.1	1.3E-2	1.6E1	
Oxidative signaling pathway	27	27	1.0	1.3E-2	1.6E1	
Estrogen signaling pathway	25	25	0.8	1.9E-2	2.3E1	
Adrenergic signaling in cardiomyocytes	34	34	1.1	1.9E-2	2.3E1	
Dilated cardiomyopathy	22	22	0.7	2.0E-2	2.3E1	
Cholinergic synapse	27	27	0.9	2.3E-2	2.7E1	
Aldosterone synthesis and secretion	21	21	0.7	2.6E-2	2.9E1	
ECM-receptor interaction	22	22	0.7	2.9E-2	3.2E1	

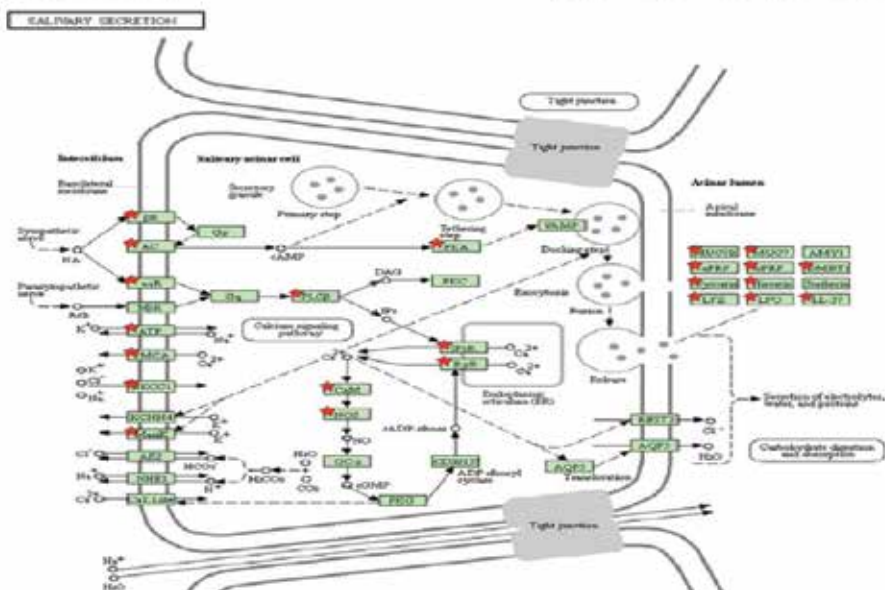


Figure 7. Pathway analysis of salivary proteins using DAVID® resulted in expected output and identification of salivary secretion as the pathway with the highest number of expressed proteins. Data courtesy of Zofia Świątczak (Master Thesis).

An exemplary result of the pathway analysis of a salivary sample using DAVID[®] is shown in **Figure 7**.

Finally, the analysis of all generated data and extracted information shall enable detection of putative biomarkers for diseases and therapy monitoring.

6. Application of salivary proteomics analysis for clinical research

As already mentioned, analysis of the salivary proteome can be applied to study a large area of conditions and diseases. The most intensively studied area of saliva as a diagnostic tool was its appliance for dental [21, 57, 64, 65], oral cancer [31, 66], diabetes [67, 68] or gastric cancer [58]. Furthermore, salivary proteomics was also applied for studying neurological and psychiatric disorders [69].

6.1. Application of salivary proteomics for diagnostics of oral diseases

The use of proteomics for diagnostics and treatment of oral diseases has been described in a number of publications. Jancsik et al. [70] describe the use of salivary proteomics to identify squamous oral cancer in diabetes patients. Authors have performed an additional sample homogenization, which is rarely described in other approaches to analyze saliva. Following analysis was performed by applying 2D gel separation of proteins and MS analysis using MALDI-TOF without the previous chromatographic separation and fractionation of proteins. It is known that inflammatory processes have a well-documented carcinogenetic role. Patients suffering from type-2 diabetes have also a higher risk of inflammatory diseases in the gastrointestinal tract such as ulcerative colitis or Crohn's disease. These patients have also a higher risk of developing gastrointestinal cancer. It was shown that the incidence of developing benign tumors, leukoplakia, and malignancies was significantly increased in the group of patients with diabetes than in the healthy control group. The authors have shown a discovery of several putative biomarkers such as, e.g. Annexin A8-like, Annexin A8-like 1, Tyrosine kinase, AX969656, Protein kinase, Peroxiredoxin-2, and Annexin A2. Annexins are known to be overexpressed in colorectal cancer but also to have altered in tumorigenesis in several types of tumor. Furthermore, loss of Annexin A1 has been found to be an early event in esophageal squamous cell carcinoma. Obviously, these results show that diabetic patients have a higher risk of developing esophageal squamous cell carcinoma than the control healthy group and close monitoring shall be applied for early detection.

Delaleu et al. [71] have performed a particularly interesting and thorough investigation of the salivary proteome from patients suffering from Sjögren's syndrome. Salivary proteome was analyzed using a 187-plex capture antibody-based assay, and the salivary proteomic biomarker profiles were generated from patients with primary Sjögren's syndrome, patients with rheumatoid arthritis, and from asymptomatic controls. Authors were able to characterize putative biomarkers by detecting significant changes in 61 and 55 proteins, respectively, in samples of patients compared to that of donors without the diagnosis of Sjögren's syndrome. Authors were able to detect, based on 4-plex and 6-plex biomarker signatures, markers

including interleukin-4 (IL-4), IL-5, and clusterin. Accurate prediction of an individual's group membership was achieved for at least 94% of cases.

Winck et al. [72] analyzed the salivary proteome in order to decipher the immune response in oral cancer based on the salivary proteome and the extracellular vesicles isolated from saliva. The authors were able to identify significant differences in processes related to inflammatory and humoral immune responses, to peptidase inhibitor activity, iron coordination, and protease binding. Based on identifications achieved, the two classes of individuals (healthy versus patients with Oral Squamous Cell Carcinoma) were distinguished with 90% accuracy based solely on the proteomics data. The authors have used the label-free approach to quantify the identified proteins. Although both groups of peptides share the great majority of identified proteins, some identified proteins were present only in the healthy or only in the group diagnosed with cancer. Authors described that out of many differentially expressed proteins, only the protein peptidyl-prolyl cis-trans isomerase A (also known as cyclophilin-A) was statistically significant in the analysis of the mean survival time of patients, with reduced abundance of PPIA being a factor that may predict poor prognosis of OSCC patients.

6.2. Application of salivary proteomics for diagnostics of diabetes

Sedentary lifestyle paired with unhealthy food, environmental derogation, and stress situations have led to significant increase in diabetes patients worldwide. Early detection of biomarkers would enable more efficient therapy and possible delay of the diseases onset or even a prevention of the outbreak. Caseiro et al. [67] have described the use of proteomics to study Diabetes mellitus (DM) type-1. Authors have performed a quantitative proteomics analysis using the chemical proteomics approach and chemical labeling using iTRAQ. Here, sample from patients diagnosed with diabetes and from healthy subjects was pooled and processed prior to LC-MS/MS measurement and bioinformatics analysis of generated data. In addition to performing iTRAQ labeling and quantitation, authors have chosen for separating peptides by applying two-dimensional separation using high pH reversed phase chromatography. It is remarkable that authors also identified endogenous salivary peptides that are mostly ignored using MALDI-TOF and combined the results to identify more than 400 proteins. Authors used the data obtained to evaluate protein expression for patients with retinopathy, nephropathy, and no complications with the salivary proteome of healthy donors. Identification of the bactericidal/permeability increasing protein-like 1 (BPI) and pancreatic adenocarcinoma upregulated factor (PAUF) in the saliva of all diabetics clearly suggests that the activation of the immune system in type 1 DM is the most prominent process. One of the proteins, BPI, is an essential component of the innate immune system with bacteriostatic and bactericidal effects against gram-negative bacteria through lipopolysaccharides binding. The PAUF is an endogenous ligand of Toll-like receptor 2 (TLR2) and Toll-like receptor 4 (TLR4), and it is also involved in the inflammatory response, which seems to be more pronounced and more prominent in patients with retinopathy and nephropathy considering the high salivary levels of alpha-2-macroglobulin, defensin alpha 3 neutrophil-specific, leukocyte elastase inhibitor, matrix metalloproteinase-9, and neutrophil elastase.

As authors also performed analysis of the peptidome, interesting data were obtained that supported the hypothesis that Diabetes Mellitus (DM)-related proteins have higher susceptibility

to proteolysis and evidenced an increased content of some specific protein fragments in saliva, which have been shown to be related with bacterial attachment and the accumulation of phosphopeptides. Especially, the accumulation of phosphopeptides seems to be involved in tooth protection against erosion and the level of their expression and phosphorylation might be a measure for dental damage that can be sustained in diabetes patients. The proteolytic fragments from bPRP1, bPRP2, and aPRP, in particular, might be considered for monitoring the disease pathogenesis with potential use for as early detection markers.

Rao et al. [73] applied multidimensional HPLC-MS/MS proteomics analysis for investigation of salivary proteins in patients with Diabetes Mellitus type-2. A Strong Cation Exchange separation of tryptic peptides from saliva of diabetic, prediabetic, and healthy subjects was performed prior to reversed phase separation and MS/MS detection. More than 400 proteins were identified and characterized and label-free quantitation was applied. As with other analysis, proteins involved with metabolic and inflammatory processes were detected in the saliva of subjects with diabetes. An important finding of the study was achieved in the prediabetic saliva: Salivary biomarkers of established Diabetes Mellitus were identified by proteomic profiling to be also differentially abundant in the saliva of patients with impaired glucose tolerance (IGT) alone and IGT + IFG (impaired fructose tolerance). These results were further confirmed by direct Western immunoblot and ELISA analyses.

The authors showed that the relative increase of some of these putative markers is associated with progression of prediabetes to the diabetic state. Therefore, systematic analysis of these putative biomarkers in prediabetic saliva, as well as their variability in individual samples, by immunoassays is of extreme importance for early acting and treatment of patients, which can prevent cardiovascular complications and mortality in diabetes patients.

An important aspect of patients suffering from DM type-2 is the severe retinopathy that can lead to blindness. Chee et al. [74] performed quantitative proteomics analysis of salivary proteins from patients diagnosed with DM type-2 without retinopathy, which served as controls and patients with DM type-2 and retinopathy. Quantitative proteomics analysis was performed by applying iTRAQ labeling and peptide separation before MS/MS detection was performed on a 50-cm nano column. Authors identified more than 300 proteins but have selected only the fully labeled pairs for quantitative analysis, a total of 119 proteins. Authors identified that increased proteins were predicted to be defense proteins and metabolic proteins suggesting that the expression of salivary defense and metabolic proteins is related to diabetic retinopathy. These results confirmed the report by Fernandez-Real et al. [75] that defense response proteins were elevated in type-2 diabetic patients and this gradually led to surging of metabolic proteins.

6.3. Application of salivary proteomics for psychiatric and neurological diseases

Fields of neurology and psychiatry urgently need new biomarkers for objective and earlier diagnoses of conditions attributed to the central nervous system (CNS). Proteomics and other "omics" technologies are being increasingly applied for these discoveries. Henskens et al. [76] performed analysis of salivary proteins already in 1996 for patients treated with different medications for epilepsy. This work was not performed by using nowadays technology, however, several salivary proteins in saliva from epileptic patients, who were medicated with

different antiepileptic drugs (namely phenytoin, valproate, and carbamazepine), were found to be increased and were compared with protein levels in the saliva of healthy control subjects. It was also found that, for all patient groups, the specific amylase activity was increased up to twofold. On the other side, absolute and relative concentrations of cystatin S were diminished in all samples, but particularly strong in patients using either valproate or phenytoin. These data suggest that use of antiepileptic drugs over long periods may cause a decrease of salivary proteins such as sIgA and cystatins, which are involved in the protection of the oral cavity against microbial infections and, therefore, these patients suffer more complications related to gingiva and oral cavity in general.

Ngounou Wetie et al. [77] have investigated the use of salivary proteins as possible markers for early onset of Autism Spectrum Disorders (ASDs). Authors have identified increased levels of apolipoproteins apoA1 and apoA4 and of serum paraoxonase/arylesterase 1 (PON1) in ASD sera compared to healthy controls in blood serum and have tested the hypothesis that levels of these peptides might also be elevated in saliva. Authors found statistically significant differences in expression of a number of salivary proteins such as elevated prolactin-inducible protein, lactotransferrin, Ig kappa chain C region, Ig gamma-1 chain C region, Ig lambda-2 chain C regions, neutrophil elastase, polymeric immunoglobulin receptor and deleted in malignant brain tumors 1. Identifications made support the hypothesis that immune system disturbances may be present in individuals with ASDs.

Castagnola et al. [78, 79] have applied a proteomics approach for studying the naturally occurring peptidome of human saliva in children diagnosed with ASDs. The study revealed that naturally occurring peptides in the saliva of children with ASD can bear multiple phosphorylations. The phosphorylation level of four specific salivary phosphopeptides, identified in this study, statherin, histatin 1, and acidic proline-rich proteins for both entire and truncated isoforms, was found to be significantly lower in autistic patients, with hypophosphorylation of at least one peptide observed in 18 ASD subjects (66%). Authors suggest that different phosphorylation and hypophosphorylation of salivary peptides suggest potential asynchronies in the phosphorylation of other secretory proteins. These proteins could be relevant in the development of central nervous system during embryonic development or in early infancy. Furthermore, obtained results suggest that naturally occurring salivary phosphopeptides might help to detect and discriminate a subgroup of ASD patients.

6.4. Application of salivary proteomics for dentistry

Saliva has a continuous and intensive contact and interaction with human teeth and plays an important role in cleaning the tooth surface and antimicrobial defense. Salivary proteomics and its role in dentistry have been studied in a number of experiments. Some of these have addressed the role of salivary proteins in edentulous patients diagnosed with DM. Byrd et al. and Border et al. [80, 81] have addressed the problem of edentulous patients with Denture stomatitis (DS) and DM type-2.

Denture stomatitis refers to an inflammatory condition of the mucosal tissue underneath the denture, which could lead to severe health problems. Clinical classification of DS distinguishes

three types: type 1 (DS I), type 2 (DS II), and type 3 (DS III), referring to clinically localized mild, localized moderate, and generalized tissue inflammation [80]. Authors have performed a quantitative proteomics analysis based on label-free quantitation and using two different MS platforms—an Orbitrap instrument and a Time-of-flight instrument. Interestingly, proteins were detected as differentially expressed between the two LC/MS systems. Protein expression was also different depending on the severity of DS. Authors have observed different levels of protein expression between different stages of DS and between the number of identified and quantified proteins for different disease stages and have identified serum proteins in the saliva of patients with DS III, e.g. ceruloplasmin, hemoglobins, serotransferrin, and albumin, which suggest that DS III patients experience higher level of inflammation and protein leakage from blood into saliva.

An interesting approach has been undertaken by Kaczor-Urbanowicz et al. [82] to study tooth absorption during an inflammatory process caused by orthodontic tooth movement. The orthodontically induced inflammatory root resorption (OIIRR) occurs as a consequence (the most prevalent and unavoidable) of orthodontic tooth movement. Authors have applied 2D-gel separation for salivary proteins upon depletion of abundant proteins (amylase, serum albumin, and IgG), and separated the tryptic peptides using HPLC followed by MS/MS detection. Identified proteins were quantified using the label-free approach. Authors were able to identify more than 700 proteins, which were revealed by quantitative MS of which different numbers were identified for different groups of patients. The strength of this study lays with the depth of the analyzed proteome and the significance of the results. Authors performed the bioinformatic analysis for proteins, which were found to be more than threefold increased. Different patient groups revealed different results although they all have been diagnosed with the tooth resorption. The moderate-to-severe root resorption young group revealed 38 functional clusters associated with acute and dynamic processes and in the moderate-to-severe root resorption adult group, other 16 functional clusters were found and those were related to less dynamic and slower processes. For the young group, these processes included the regulation of acute inflammatory response, defense response, response to stress, response to wounding or healing as opposed to apoptosis, glycoproteins expression, cell adhesion, signal peptides, etc. in the adult group with moderate-to-severe processes. Finally, a number of new putative biomarkers were identified, and these might be used to produce a clinical test that would serve along with radiography to perform a fast and more reliable diagnosis.

7. Conclusion

In summary, salivary proteomics is an upcoming approach for both basic and clinical research with a significant potential for use in fast diagnostic approaches. Not only the analysis of salivary proteins but also the analysis of endogenous peptides in saliva and their posttranslational modifications shall be addressed and targeted. Although a number of studies have been published and more are to come, more research is required to validate the discovered putative biomarkers so far.

Current proteomics approach for the analysis of oral fluid is not yet suited for daily routine in clinical diagnostics. However, it can help discovering biomarkers for which immunological tests such as ELISA can be developed.

Furthermore, it is essential to develop and curate a comprehensive database for the salivary proteome and establish standard conditions for sample collection and processing until the MS analysis.

The space was scarce in this chapter to address more of the clinical approach of salivary proteomics, but the researcher is encouraged to stay focused and follow the further development.

Current development shows that, without any doubt, this process will continue and will yield more biomarker candidates in the future.

Author details

Goran Mitulović

Address all correspondence to: goran.mitulovic@meduniwien.ac.at

Medical University of Vienna, Clinical Department of Laboratory Medicine, Vienna, Austria

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Approach to Diagnosis of Salivary Gland Disease from Nuclear Medicine Images

Michihiro Nakayama, Atsutaka Okizaki,
Kaori Nakajima and Koji Takahashi

Additional information is available at the end of the chapter

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Abstract

Nuclear medicine images can help in the diagnosis and assessment of some salivary disorders. $^{99m}\text{TcO}_4^-$, gallium-67-citrate scintigraphy will be an indication of the function of salivary gland together and it will be used for the diffuse diseases such as sialadenitis, Sjögren's syndrome, sarcoidosis, glossopharyngeal paralysis, and irradiation. It is also effective for distinguishing benign tumor lesion with Warthin's tumor and others. Moreover, fluorodeoxyglucose positron emission tomography (FDG-PET) is an indispensable modality for determining the localization, focal lesions, and staging of many malignant tumors, the fluorodeoxyglucose (FDG) accumulation is visually and semi-quantitatively assessed using the standardized uptake value (SUV), which is the ratio of uptake to the injected dose per unit body weight. Also for radioactive iodine therapy, attention should be paid to adverse reactions. It is important to note that acute/chronic salivary gland disorders are associated with radioiodine therapy for the treatment of postoperative thyroid cancer. Coordination among healthcare providers including nurses, radiological technologists, and doctors of all departments involved in treatment is important for achieving effective outcomes.

Keywords: scintigraphy, $^{99m}\text{TcO}_4^-$, SPECT, FDG-PET, salivary gland disorders

1. Introduction

Radionuclide imaging, commonly used for the diagnosis of salivary gland diseases, consists of salivary scintigraphy using $^{99m}\text{TcO}_4^-$, gallium-67-citrate (^{67}Ga) scintigraphy in inflammation, and fluorodeoxyglucose positron emission tomography (FDG-PET). It is important to note that acute/chronic salivary gland disorders are associated with radioiodine therapy for the treatment of postoperative thyroid cancer.

2. Salivary gland scintigraphy

2.1. Mechanism of uptake

Salivary glands secrete saliva. The parotid, submandibular, and sublingual glands are called the “major salivary glands.” Salivary epithelial cells have a sodium/iodide symporter (NIS), which takes up univalent anions such as Cl^- and I^- and concentrates them (**Figure 1**). The concentrated anion is secreted into saliva. Administered $^{99\text{m}}\text{TcO}_4^-$ (**Figure 2**) is taken up by the salivary glands through NIS, similar to Cl^- . Thus, intravenously administered $^{99\text{m}}\text{TcO}_4^-$ accumulates mainly in the parotid and submandibular glands and is excreted into saliva [1]. After the accumulation of the radionuclide, loading of citric acid, such as lemon juice, stimulates the secretion of saliva, which indicates salivary gland function. Salivary gland scintigraphy is useful for differentiating salivary gland tumors because Warthin’s tumors and oncocytomas, which are benign, retain $^{99\text{m}}\text{TcO}_4^-$.

2.2. Testing procedure

Because salivary gland function is affected by food intake, 1 hour of fasting is needed before testing.

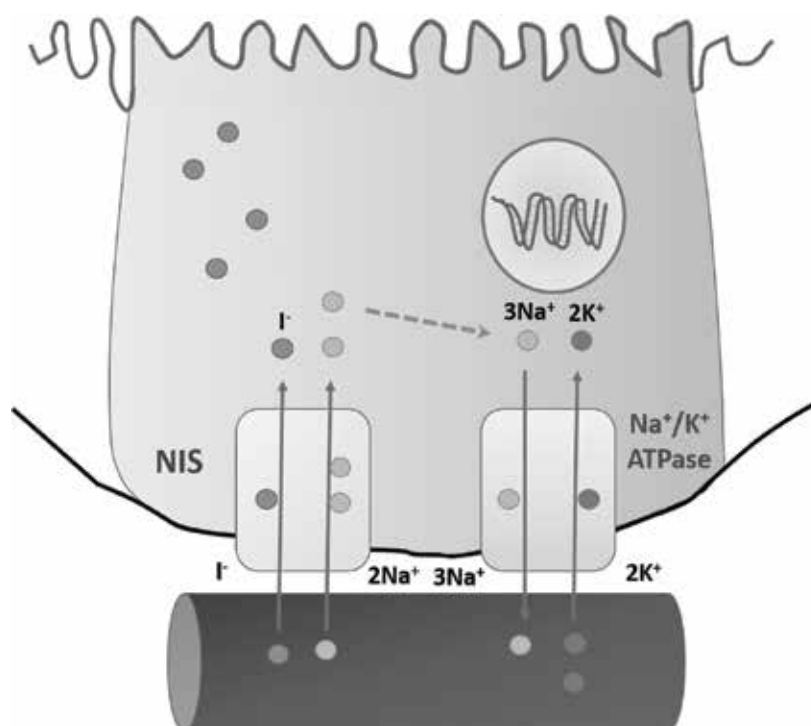


Figure 1. Iodide uptake mechanism of sodium/iodide symporter. Sodium/iodide symporter transports two sodium ions and one iodide ion into the cytoplasm together.

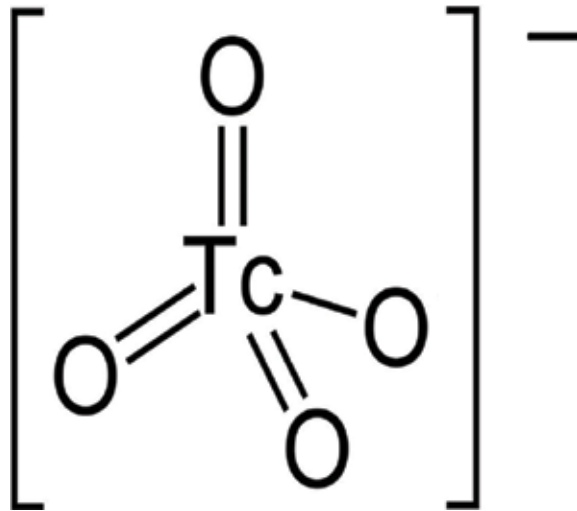


Figure 2. Structural formula of $^{99m}\text{TcO}_4^-$. $^{99m}\text{TcO}_4^-$ is incorporated into the salivary glands through NIS, similar to Cl^- .

For the kinetic analysis of salivary gland function, 185–370 MBq (5–10 mCi) of $^{99m}\text{TcO}_4^-$ is intravenously administered. A dynamic scan (anteroposterior view) is performed at 5-minute intervals for 30 minutes. The thyroid gland should be included in the area. Citric acid (e.g., lemon juice) is instilled into the oral cavity 20 minutes after the intravenous injection to stimulate the secretion of saliva. The regions of interest are set at the bilateral parotid and submandibular glands, and at background regions to generate time-activity curves (TAC). The TAC is used to determine the function of individual salivary gland. For the diagnosis of tumors and morphology of salivary glands, the intravenously administered dose is 370–740 MBq, and the anteroposterior and lateral images are obtained 10–15 minutes after intravenous administration (**Figure 3**). When the assessment of the tumor is difficult due to the physiologic uptake in the normal salivary glands, washout by stimulating the secretion of saliva is useful.

2.3. Imaging evaluation

2.3.1. Normal images

Salivary gland scintigraphy provides information about the morphology and function of the salivary glands and the procedure is easier than that of sialography.

2.3.2. Dynamic scans and TAC

The uptake in the bilateral parotid and submandibular glands begins less than 1 minute after intravenous administration of the radionuclide and increases over time. The uptake in the parotid glands is equal to or greater than that in the submandibular glands. The sublingual glands are not visible, though the reason is unknown. After the stimulation of saliva secretion, the uptake rapidly declines in all four glands and subsequently rises again. The percentage

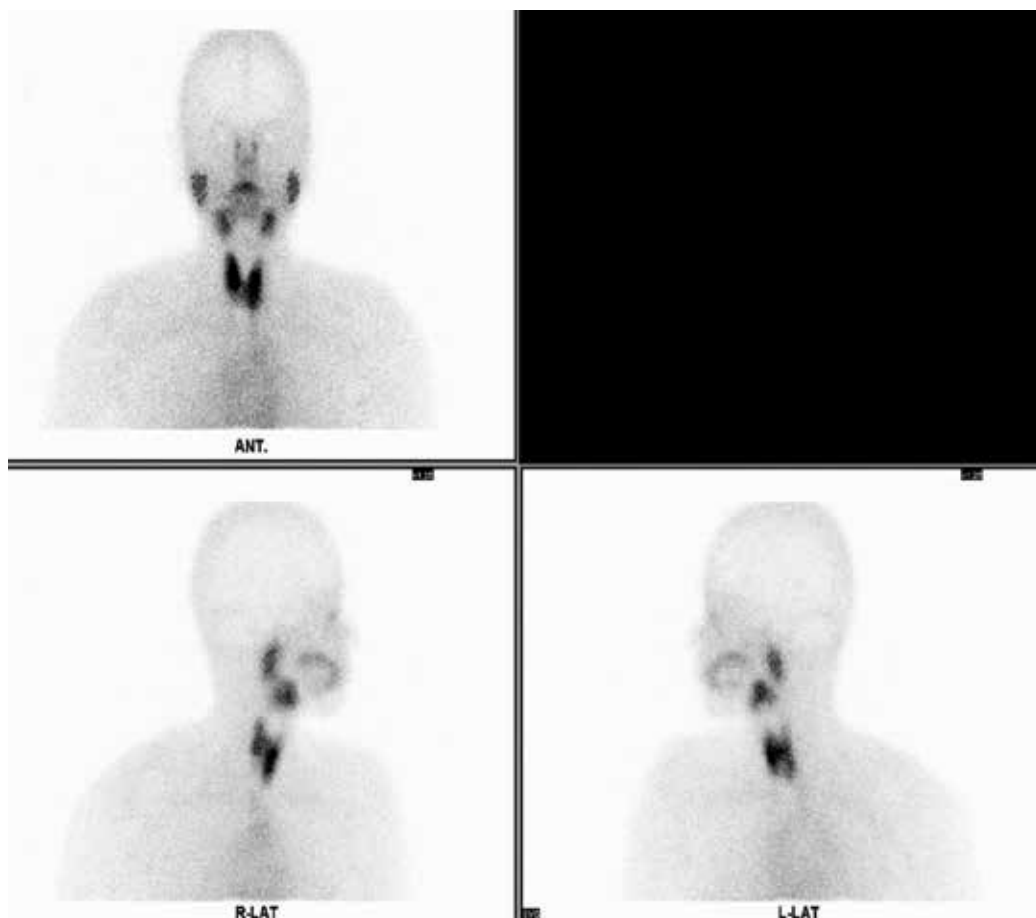


Figure 3. Static images in normal case. The radionuclide uptake of the parotid and submandibular glands is equal to or lower than that of the normal thyroid gland.

of washout is calculated using the counts at the maximum uptake and those at the minimum uptake seen after the stimulation of saliva secretion in each gland. The washout (%) is 50% or higher in the normal salivary gland (**Figure 4**).

2.3.3. *Static images*

The radionuclide uptake of the parotid and submandibular glands are equal to or lower than that of the normal thyroid gland, and mild uptake appears in the nasal and oral cavities. On the lateral view, the parotid gland is clearly shown, but the submandibular gland overlaps with the contralateral submandibular gland.

2.4. Diagnosis of salivary gland tumors

Warthin's tumors and oncocytomas are derived from the epithelial cells of excretory ducts and do not communicate via excretory ducts (**Figure 5**). Thus, $^{99m}\text{TcO}_4^-$ is taken up by the

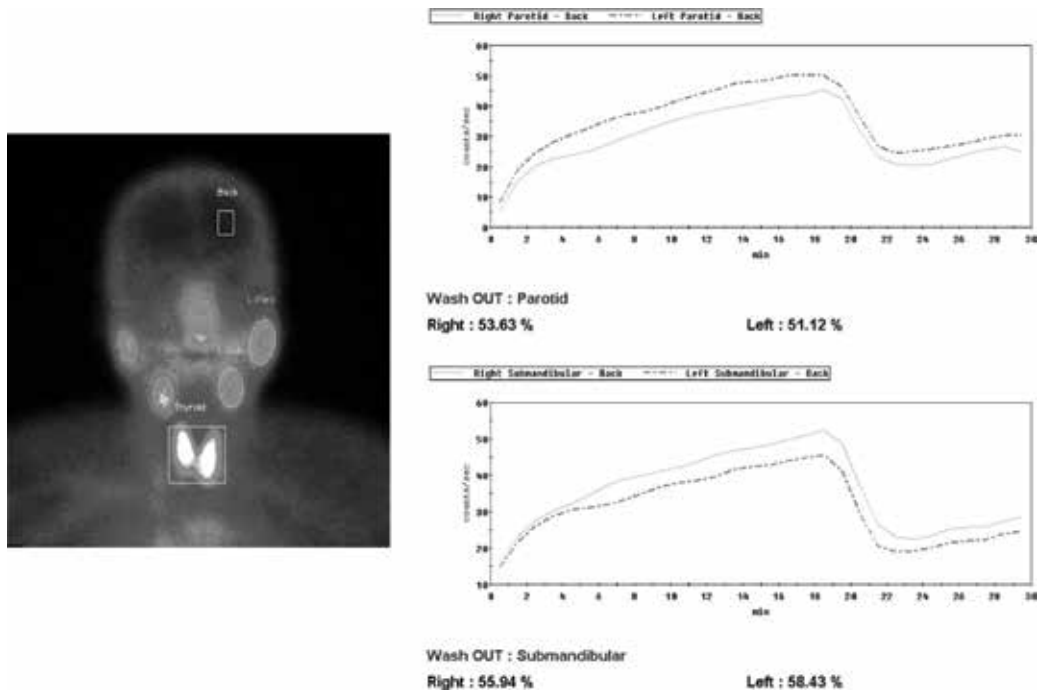


Figure 4. TAC of the normal salivary gland. After intravenous administration of the radionuclide, uptake in the parotid and submandibular glands increases over time. At 20 minutes after administration, saliva secretion is stimulated. Counts in the salivary glands rapidly decline and then gradually increase again.

solid component of these tumors and is not eliminated after the stimulation of saliva secretion. However, the uptake may be reduced in Warthin’s tumor mainly with a cystic component. The diagnostic accuracy of salivary gland scintigraphy for Warthin’s tumors and oncocytomas is around 90%, but these cannot be differentiated in salivary gland scintigraphy. Because Warthin’s tumors develop bilaterally in 5–20% of cases, the contralateral parotid gland should be carefully evaluated. Meanwhile, because $^{99m}\text{TcO}_4^-$ is not taken up by pleomorphic adenomas or parotid gland cancer (defect images), salivary gland scintigraphy is useful for differentiation [2].

2.5. Kinetic analysis of salivary gland function

The indications for kinetic analysis of salivary gland function include Sjögren’s syndrome, acute/chronic sialadenitis, and facial/glossopharyngeal nerve palsy. Salivary gland function is assessed based on dynamic images and TAC. In general, chronic sialadenitis shows a decreased uptake while acute sialadenitis shows an increased uptake; acute/chronic sialadenitis shows a reduced or no response to stimulation of saliva secretion. Uptake in salivary glands is barely evident in patients with severe Sjögren’s syndrome (Figures 6 and 7). The severity of the reduction in washout after the stimulation of saliva secretion well correlates with the results of the Saxon test. Thus, kinetic analysis reflects the severity of Sjögren’s syndrome, allowing an objective assessment of salivary gland function [3].

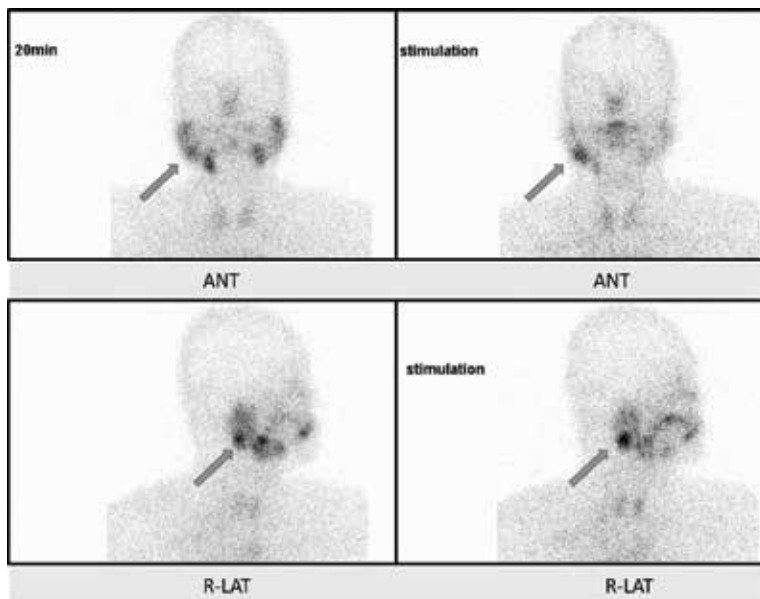


Figure 5. Warthin's tumor. On the image 20 minutes after administration, the radionuclide is taken up by the right parotid gland. After the stimulation of saliva secretion, the radionuclide remains in the right parotid tumor. The diagnosis was Warthin's tumor.

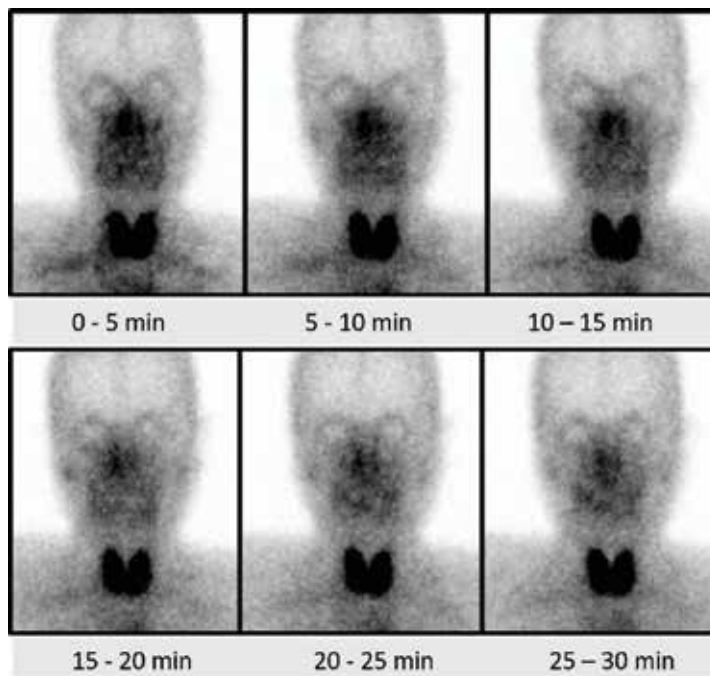


Figure 6. Sjögren's syndrome. All four salivary glands show decreased uptake.

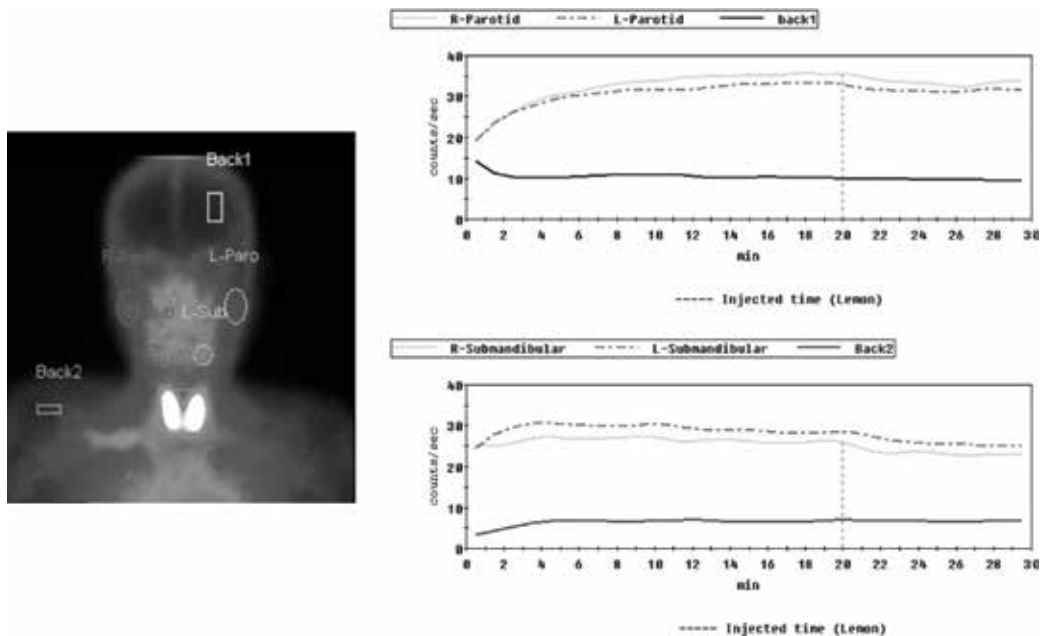


Figure 7. TAC of the Sjögren's syndrome. All four salivary glands show decreased uptake and a poor response to stimulation of saliva secretion 20 minutes after administration. TAC is useful for assessment.

3. Scintigraphy of inflammation (^{67}Ga scintigraphy)

3.1. Mechanism of uptake

^{67}Ga administered intravenously binds to transferrin, a serum protein, and is transported into cells through transferrin receptors. The carbon atom of citrate stabilizes the bond between ^{67}Ga and transferrin. Transferrin receptors that bind to ^{67}Ga distributed in lysosomes and cytoplasm are often present in tumor and inflammatory cells, which show intense uptake of ^{67}Ga .

3.2. Testing procedure and imaging evaluation

^{67}Ga is intravenously administered at a dose of 185–555 MBq. Imaging is performed 48–72 hours after intravenous administration to visualize the distribution of the radionuclide. ^{67}Ga is excreted from the kidney and intestinal tract within 24 hours after administration and is mainly excreted by the liver. Intense uptake of ^{67}Ga is noted in the liver, bone, and spleen 48–72 hours after administration. ^{67}Ga is known to be taken up by inflammation and tumors; however, the sensitivity of ^{67}Ga scintigraphy is low for malignant tumors, while the negative predictive value is high. Thus, a negative finding of focal uptake is likely to represent a benign lesion or low-grade tumor. Focal uptake in the parotid gland on ^{67}Ga scintigraphy is useful for the supplemental diagnosis of Warthin's tumor. Meanwhile, with increased diffuse bilateral uptake, differential diagnosis includes sarcoidosis (**Figure 8**),

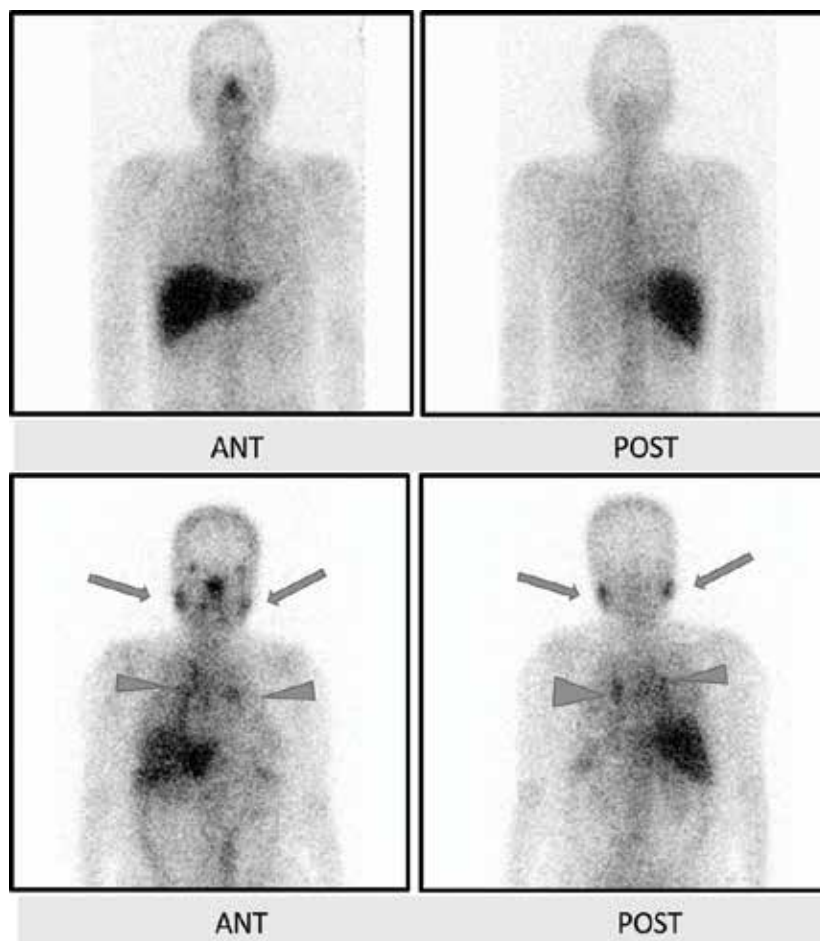


Figure 8. ^{67}Ga scintigraphy. Upper column, normal image; lower column, sarcoidosis. Uptake is noted in the bilateral parotid glands (arrow), and mediastinal/hilar lymph nodes (arrowhead).

IgG4-related disease, Sjögren’s syndrome, and Mikulicz disease. In recent years, ^{67}Ga scintigraphy for tumor diagnosis has been increasingly replaced by FDG-PET, as described below.

4. PET

4.1. Mechanism of uptake

FDG-PET is a critical modality for determining the localization, focal lesions, and staging of many malignant tumors, as well as for their follow-up observation. It is also essential for the clinical management of salivary tumors [4–8]. Like glucose, fluorodeoxyglucose (FDG)

is taken up by cells via glucose transporters and phosphorylated; however, unlike glucose, FDG remains in cells after phosphorylation. In general, glucose transporters and glucose metabolism are increased in tumor cells, leading to an increased uptake of FDG (Figure 9). The widespread use of PET combined with computed tomography (PET/CT) has increased the diagnostic accuracy by compensating for PET disadvantages, including poor spatial resolution and lack of anatomic information. Moreover, PET combined with magnetic resonance imaging (PET/MRI) has recently emerged.

4.2. Testing procedure

Fasting is required for 4 hours before testing and intake of liquids with sugar content is prohibited. FDG (Figure 10) is intravenously administered at a dose of 185–444 MBq (5–12 mCi). Imaging is performed 60–90 minutes after administration to visualize distribution. The accumulation is visually and semi-quantitatively assessed using the standardized uptake value (SUV), which is the ratio of uptake to the injected dose per unit body weight.

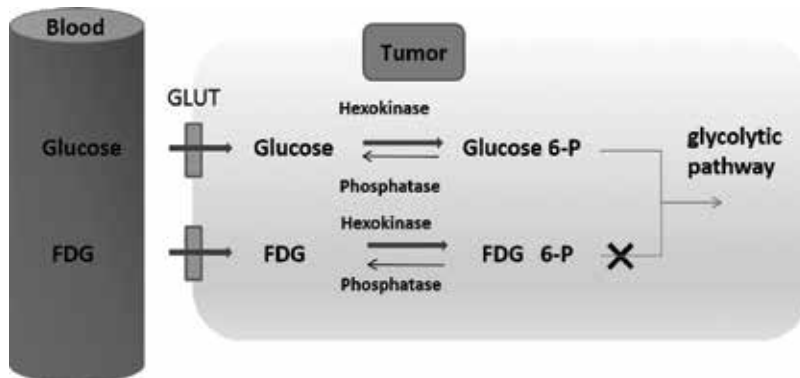


Figure 9. Mechanism of FDG uptake. Like glucose, FDG is taken up by cells via glucose transporters and phosphorylated; however, unlike glucose, FDG remains in cells after phosphorylation.

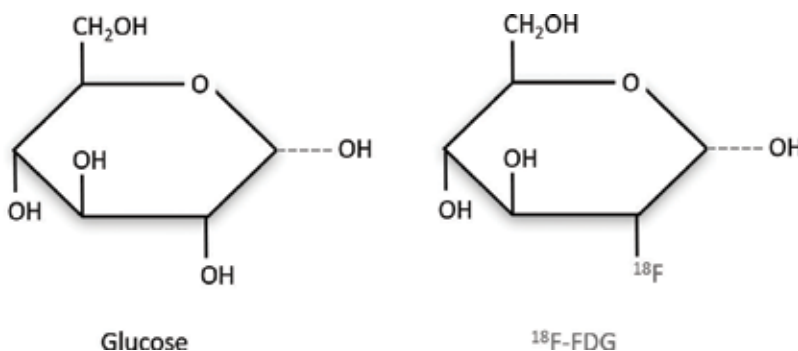


Figure 10. The chemical structure of FDG. The chemical structure of FDG is identical to that of 18F (one of the hydroxy groups of glucose that is replaced by a positron-emitting radionuclide).

4.3. Normal uptake

In the head and neck areas, many structures show physiologic uptake, including salivary glands, nasal and sinonasal mucosa, extraocular muscles, and lymphoid tissue. Because artifacts due to dentures are also often seen, information on CT or MRI images is useful [8].

4.4. Diagnosis of salivary gland tumors

Some lesions are difficult to differentiate from normal structures, postoperative changes, and inflammatory changes on CT or MRI images alone. However, those lesions can be diagnosed through the combined use of FDG-PET [4, 5, 7]. FDG-PET for medical evaluation or staging of a malignant tumor may incidentally reveal a salivary gland tumor [7, 9].

The differentiation of benign from malignant parotid tumors is difficult based on the results of FDG uptake alone. Moreover, the differentiation of benign from malignant salivary tumors is often difficult based on the comparison of the results of SUVmax alone. A malignant tumor tends to show more intense FDG uptake than a benign tumor; however, benign tumors, such as Warthin's tumor and pleomorphic adenoma (**Figure 11**) [10–12], also show high FDG

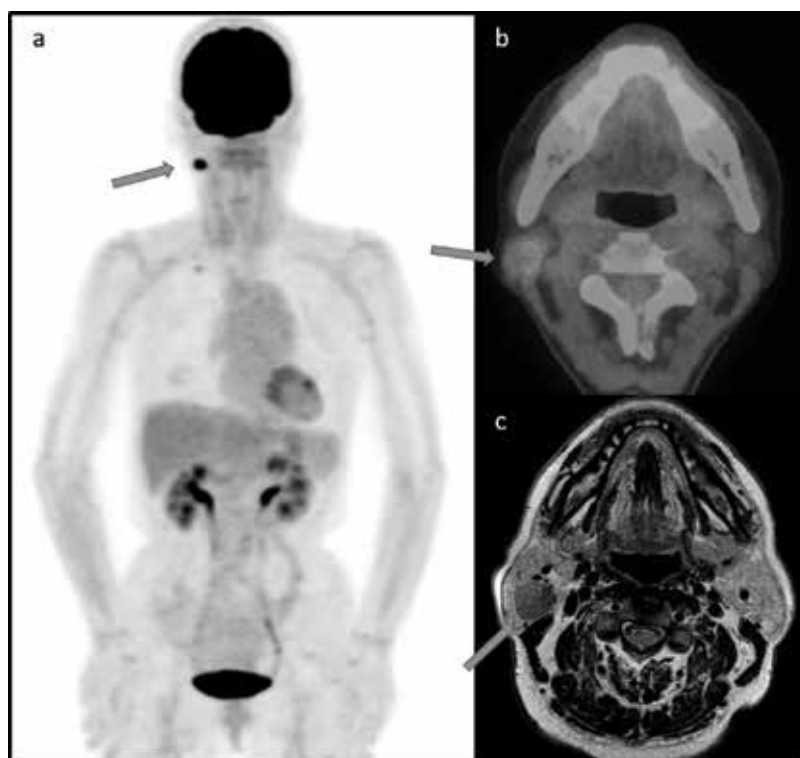


Figure 11. Warthin's tumor. (a). PET MIP image. (b). PET/CT fusion image. (c). T1-weighted MRI image. FDG is strongly taken up by the tumor (SUVmax: 9.4) in the right parotid gland. MRI also provides many findings consistent with malignancy. Salivary gland scintigraphy is useful for differentiation.

uptake. Some studies have reported that the differentiation of benign from malignant salivary tumors is possible with the use of indices such as dual-time-point (DTP) imaging and tumor volume, in addition to SUVmax [4, 5].

The sensitivity of PET/CT is approximately 80% for the detection of preoperative primary lesions (**Figure 12**), but the accuracy for staging may vary. In particular, FDG-PET images often show false-positive results for the diagnosis of cervical lymph nodes.

Sialadenitis may show diffuse, increased uptake. However, some cases may show unilateral uptake due to the distribution of inflammation and may be difficult to differentiate from a tumor.

FDG-PET, which provides information about systemic metabolism, is very useful for detecting primary or recurrent lesions for the determination of treatment strategy in cases with highly-malignant tumors requiring aggressive treatment. FDG-PET aids in detection of local involvement, regional lymph node metastasis, distant metastasis, and dissemination for the clinical staging and restaging. It is also useful in the detection of an incidental

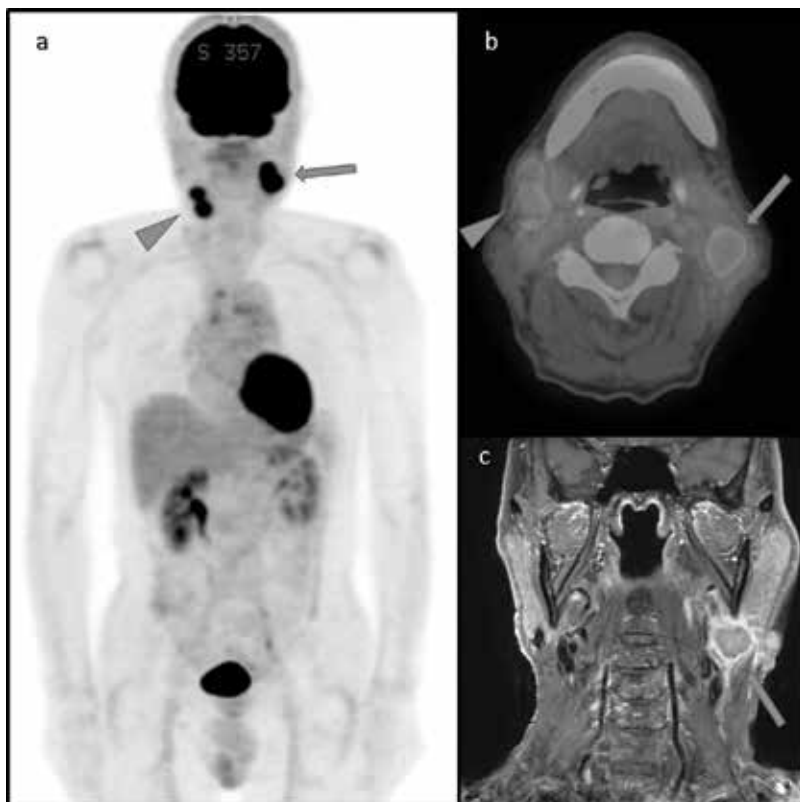


Figure 12. (a). PET MIP image. (b). PET/CT fusion image. (c). T1-weighted MRI using gadolinium-based contrast (GdT1) image. Very strong uptake of FDG (SUVmax: 15.8, arrow) is seen in the region corresponding to the left parotid tumor on the MRI image. Uptake is seen in the region of the right internal jugular lymph nodes (SUVmax: 10.2, arrow head). Parotid cancer or cervical lymph node metastasis is suspected.

cancer. Therefore, this imaging modality is essential before the initiation of treatment and for patient follow-up [4, 5, 13–15].

5. Salivary gland disorders associated with radioiodine therapy

Radioactive iodine (RAI) therapy is the most widely used treatment for differentiated thyroid cancer and has a long history. Because NIS is expressed in the salivary glands, ^{131}I is also incorporated into the salivary glands. Not only thyroid cancer but also salivary glands are irradiated; and thus, acute/chronic sialadenitis may develop.

5.1. Acute salivary gland disorders

Sialadenitis is one of the most common adverse reactions to RAI therapy. Acute-phase sialadenitis causes swelling of the major salivary glands and pain (especially while eating) within one to a few days after oral administration of ^{131}I . Although sialadenitis ameliorates without treatment, xerostomia may develop during the chronic phase and significantly impair the quality of life. Thus, when symptoms develop, sialadenitis should be treated immediately [16, 17]. The parotid glands may be affected more often than the submandibular glands. Steroids are more effective than nonsteroidal anti-inflammatory drugs. Cooling and frequent rinsing with cold water may relieve symptoms. Taste dysfunction is characterized by reduced acuity for salt taste and subsequently sweet taste; bitter taste remains unaffected. Taste dysfunction may become obvious later (2–4 weeks after ^{131}I administration), rather than immediately after administration. To prevent an acute-phase disorder, snacks such as lemon candy to stimulate saliva secretion may be helpful because they also promote the excretion of ^{131}I . Apitherapy using honey products is reportedly useful for preventing salivary gland disorders. Massage of the salivary glands and aroma therapy are also reported to relieve symptoms [18].

5.2. Chronic salivary gland disorders

Xerostomia may develop during the chronic phase, even in patients without acute impairment. Permanent dysfunction is reported to develop more frequently in the submandibular glands than in the parotid glands. Salivary gland scintigraphy shows a pattern of obstructive dysfunction before revealing parenchymal damage of the salivary gland. Therefore, salivary gland function may improve if salivary gland dysfunction can be identified within the period showing an obstructive pattern [19]. Salivary gland function should be monitored regularly with scintigraphy or other studies [20].

6. Conclusions

Radionuclide imaging of salivary glands has been used not only for functional assessment but also for comprehensive diagnosis including morphological information with the advent of single-photon emission computed tomography (SPECT)/CT and PET/CT. Although scientific

evidence is limited, the advantages of radionuclide scanning should be determined. For RAI therapy, attention should be paid to adverse reactions. Coordination among healthcare providers including nurses, radiological technologists, and doctors of all departments involved in treatment is important for achieving effective outcomes.

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Author details

Michihiro Nakayama*, Atsutaka Okizaki, Kaori Nakajima and Koji Takahashi

*Address all correspondence to: m-naka@asahikawa-med.ac.jp

Department of Radiology, Asahikawa Medical University, Japan

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New Approaches in the Treatment of Salivary Gland Diseases

Salivary Effects of Facial Vibrotactile Stimulation in Patients with Sjogren's Syndrome and Poor Salivation

Koichiro Ueda, Kanako Gora, Masaru Yamaoka,
Takako Sato, Kimiko Abe, Enri Nakayama,
Mituyasu Sato, Yumiko Tunoda, Sumiko Akatuka,
Sayaka Fukui, Akinari Hayashi, Teruyuki Hirai,
Sayako Ohnishi, Mayumi Hayata and Hisao Hiraba

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Abstract

We examined the effect of vibrotactile apparatus in patients with Sjögren's syndrome and others with reduced salivation in comparison to normal subjects. The most effective salivation in normal subjects was produced by 89 Hz vibrotactile stimulation with 9.8 μm amplitude on the parotid or submandibular glands vibrotactile stimuli. First, we examined by measuring the weight of dental cotton rolls positioned at the opening of the secretory duct for total salivation 3 min during resting, and then after 5-min intervals, the weights were measured every 3 min of vibrotactile stimulation on salivary glands. Furthermore, we measured facial temperature around vibrators after 2 min of vibration. We investigated 10 poor salivation patients with Sjögren's syndrome (8 patients) defined by examinations (contrast study or scintigraphic test) and others (2 patients). About 50% of patients with poor salivation gained recognition for good results, although they had periods of short-term (3 months) and long-term effects (6–7 years) during recuperation. Furthermore, facial skin temperatures on both sides of parotid glands were decreased in Sjögren's syndrome after vibration, although their temperatures were increased following recovery. Although the mechanism is not clear, we think that vibrotactile stimulation gives activation to salivary glands under the rising facial temperature.

Keywords: vibrotactile stimulation, facial skin temperature, parotid and submandibular glands, poor salivation, Sjögren's syndrome

1. Introduction

When we treat patients with reduced salivation (hyposalivation), we provide treatments such as artificial saliva, humectant, massage on the salivary glands, and so on [1]. However, treatment *with* the artificial saliva and humectant is the symptomatic treatment, and patients with handicaps *experience difficulties when they do massage*. We previously reported about *the relationship between facial vibratory effects in normal subjects and promotion of salivation*. We *performed this method* for facial vibratory effects on Sjögren's syndrome patients with poor salivation. We focused on *increase of salivation with the use of facial vibrotactile stimulation*, as reported by Hiraba et al. [2, 9, 10].

When patients continuously utilize the apparatus in future *the decrease or increase of salivation is examined from this result* [3]. In this experiment, it was necessary to make a comparison between the resting and *stimulated* salivations and to investigate the most effective frequency for increasing the salivary secretion. We examined the amount of salivation during vibrotactile stimuli with one vibrating motor (1.9 μm amplitude) on the bilateral masseter muscle belly (on the parotid glands), and in patients of Sjögren's syndrome, we asked twice practice during 15 min of morning and night. Furthermore, *the amount of salivation was explored by using a dental cotton roll positioned at the opening of the secretory duct for 3 min*. After this experiment was performed, *we made a comparison between the resting and stimulated salivations and investigated the most effective frequency for increasing salivary secretion*. When we examined normal subjects, *the effect of the increased salivation determined the difference between the resting and stimulated salivations*. We think that *total salivation after the resting phase shows conditions of day-to-day salivation, and they after the stimulating are effects of vibrotactile stimulation*.

We defined 5-min intervals as the recovery time between the resting and *stimulated* salivations from the previous pre-examinations. Furthermore, we examined temperature effects on patients with poor salivations (affected by Sjögren's syndrome) and others by the use of facial vibrotactile stimuli. Increased facial temperature by the vibrotactile stimulation showed changes of metabolism around facial skins. We will discuss the effects of vibrotactile stimulation based on these results.

2. Material and methods

2.1. Vibrotactile stimulation apparatus

The vibrotactile stimulation apparatus consists of an oscillating body and control unit, as shown in **Figure 1A** [2]. The oscillating body is composed of the headset equipped with vibrators as a substitute for positions of the bilateral microphones and vibrators utilizing the vibration electric motor (VEM) (Rekishin Japan Co., LE12AOG), as shown in **Figure 1A**. The VEM is covered in silicon rubber (polyethyl methacrylate, dental mucosa protective material, Shyofu Co.) for conglobating the parts under stimulation and preventing the warming of the VEM's temperature produced by the vibration of long periods [2]. The control unit consists

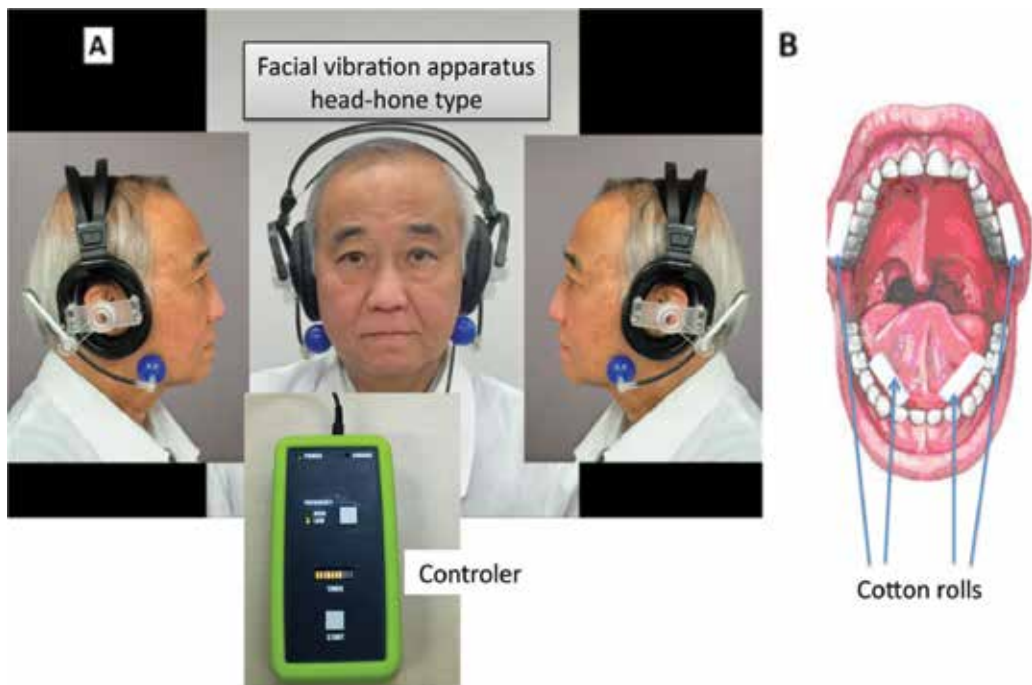


Figure 1. (A) Apparatus for facial vibration and (B) position of cotton rolls.

of three parts: the pulse width modulation (PWM) circuit, LCD monitor circuit and power supply circuit, and *interfaces* with a PWN electric motor, *delivers* vibration frequencies in the 60–182 Hz range [4].

As shown in **Figure 1**, we examined the amount of salivation during vibrotactile stimuli by two kinds of methods: on the bilateral masseter muscles belly (parotid glands) and on the bilateral parts of submandibular angle (submandibular glands). We examined the amount of salivation using a dental cotton roll (1 cm across and 3 cm length) positioned at the opening of the secretory ducts (right and left sides of parotid glands and right and left sides of submandibular and sublingual glands), during the vibrotactile stimuli on the bilateral parotid and submandibular glands, and wet cotton rolls measured for 3 min. These weights were then compared to their initiatory weights, as shown in **Figure 1B** [2].

2.2. Estimation of the stimulating salivation in normal subjects

First, we use three different frequencies, 89, 114 and 180 Hz as the vibrotactile frequency from the character of the oscillating body on the parotid glands. **Figure 1** shows the apparatus and position of rolls. To begin with, we put an exercise into practice for avoiding foreign-body sensation on the cotton rolls while setting for 3 min. Next, after 5 min of resting, we examined the amount of salivation during the 89 Hz vibrotactile stimulation for 3 min. Furthermore, after every 5 min of rest, we examined next amount of salivation during the 114 and 180 Hz vibrotactile stimuli for 3 min, respectively. We decided on 3 min for the measurement of salivation and 5 min for recovery

time from the previous experiment [2]. We carried out the examinations and used 19 normal subjects (male: 6 and female: 13, average age 22) for the resting-stimulating examination. This experiment was performed between 3 and 5 pm in a temperature-controlled room.

Second, as shown in **Figure 2**, we used three different frequencies (89, 114 and 180 Hz) and two different amplitudes (1.9 and 3.5 μm) on the parotid and/or submandibular glands. Amplitudes of vibrotactile stimuli were measured by the CCD laser displacement gauge (LK-G3000, KEYENCE Co.). After three different frequencies were attempted on the parotid glands, we explored the most effective frequency, and we arrived at a frequency of 89 Hz. We examined the frequency of 89 and 114 Hz and we used also oscillating bodies added as the frequency with double motors (one motor is 1.9 μm amplitude and double motors is 3.5 μm amplitude). Namely, the second experiment was practiced by 89 and 114 Hz with one motors (1.9 μm amplitude), and 89 and 114 Hz with double motors (3.5 μm amplitude). We examined the amount of salivation in four different trials, as shown in **Figure 2**. We carried out the examinations and used 17 normal subjects (male: 15 and female: 2, average age: 22) for the resting-stimulating examination. This experiment was performed between 3 and 5 pm in a temperature-controlled room.

Finally, as shown in **Figure 2**, since the most effective salivation by vibrotactile stimuli was at 89 Hz frequency with one motor (1.9 μm amplitude), we examined salivations on 89 Hz vibrotactile stimulation continuously for 4 or 5 days. As patients continuously utilized the apparatus, we examined if adaptation develops with everyday usage and whether or not the decrease of salivation arises. We investigated the adaptation of periods with the continuous use of vibrotactile stimuli for 4 continuous days in the same subjects. We carried out this examination and used 26

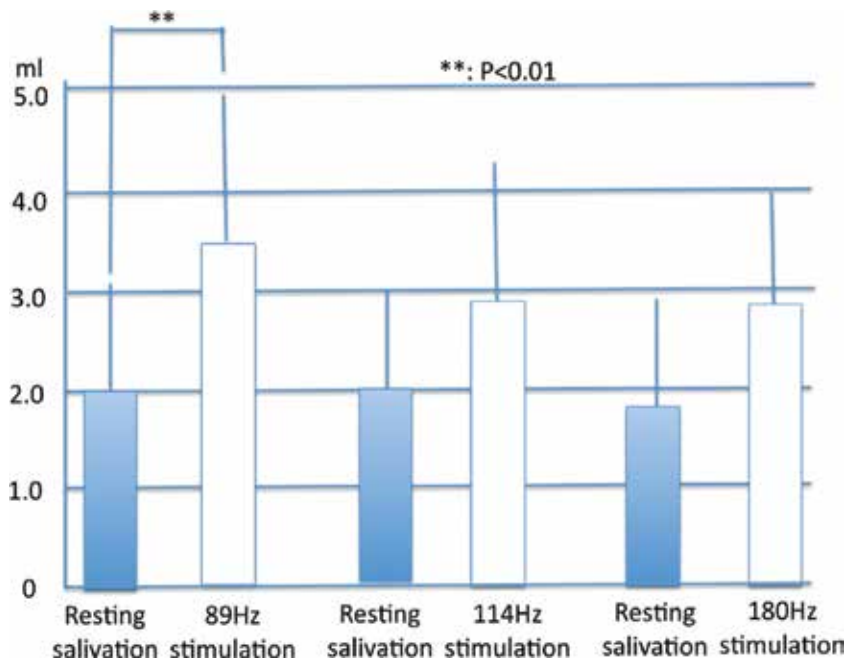


Figure 2. Salivations in each vibration frequency.

normal subjects (male: 11 and female: 15, average age 25) for the resting-stimulating examination. This experiment was performed between 3 and 5 pm in a temperature-controlled room. *In particular, we find that 89 Hz frequency and 1.9 μm amplitude is most effective for salivation.*

2.3. Vibrotactile stimuli on the parotid and submandibular glands

As shown in **Figure 1**, we examined the difference between vibrotactile stimuli on the parotid glands and the submandibular glands. First, we tried three different vibrotactile stimuli, 89, 114, and 180 Hz, on the parotid glands and explored the frequency for the most effective salivation, as shown in **Figure 2**. Next, we inferred the most effective salivation of 89 Hz with the one motor depending on vibrotactile stimuli on the parotid or submandibular glands [2]. Furthermore, we investigated the most effective salivation depending on the difference of amplitudes (1.9 and 3.5 μm amplitudes). We inferred the most effective salivation of 89 Hz with the one motor (1.9 microm amplitude) on the submandibular glands (significant difference $P < 0.05$, 89 Hz with 1.9 μm amplitude and 114 Hz with 1.9 and 3.5 μm amplitude), as shown in Hiraba et al. [2].

2.4. Total salivation after the vibrotactile stimulation on the parotid or submandibular and sublingual glands

In 89 Hz vibrotactile stimulation with 1.9 μm amplitude on the parotid glands, we observed the most effective salivation in each gland, the right parotid, left parotid, right submandibular and sublingual, and left submandibular and sublingual glands. *Vibrotactile stimuli on the parotid or submandibular glands in any case showed that at 89 Hz more effective salivation in the right and left parotid and in the left and right submandibular and sublingual glands happened in comparison with the resting salivation in each gland.* On the other hand, vibrotactile stimuli with 1.9 μm (89 and 114 Hz—1) or 3.5 μm amplitudes (89 and 114 Hz—2) on the parotid or submandibular glands were examined. The 89 Hz with one motor, was the most effective salivation in the parotid, and the submandibular and sublingual glands, and the 89 Hz with double motors, was the more effective salivation in the parotid, and submandibular and sublingual glands, as shown in Hiraba et al. [2]. From these reasons, we suggested that vibrotactile stimulation at 89 Hz with 1.9 μm amplitude showed the most effective salivation in many glands.

Finally, we assumed that 89 Hz with 1.9 μm amplitude vibrotactile stimulation *produced the most effective salivation*, and then the vibrotactile stimuli on the parotid and submandibular glands *showed* hardly any difference. We then decided to use the apparatus to patients affected by poor salivation.

2.5. Information of patients with poor salivation affected by Sjögren's syndrome and others

In **Figure 3**, we showed information of patients: eight women were *diagnosed with Sjögren's syndrome* and two women's symptoms were unexplained. *Patients with Sjögren's syndrome were diagnosed by contrast and/or scintigraphic studies. In particular, the patients with indefinite complaints were not given a definite diagnosis; nevertheless, they had poor salivation as their chief complaint.*

<u>Name</u>	<u>Definite diagnosis</u>	<u>Saligren Dose level</u>	<u>Passed years</u>	<u>Actual condition</u>
1. MZ	contrast study	dispense medication (3 drugs/day)	8 years	continued investigation
2. AS	contrast study	dispense medication (3 drugs/day)	5 years	break up for myalgia
3. NI	contrast & scintigraphic study	dispense medication (3 drugs/day)	5 years	vibration disamenity
4. KM	contrast study	dispense medication (3 drugs/day)	3 years	ope of brain cancer
5. TO	scintigraphic study	dispense medication (3 drugs/day)	6 years	contiued investigation
6. YA	contrast & scintigraphic study	dispense medication (3 drugs/day)	1.5years	cured patient
7. NA	scintigraphic study	no drug	6 years	continued investigation
8. UJ	indefinite complaint	no drug	4 years	break up for change residence
9. SG	indefinite complaint	dispense medication (3 drugs/day)	2 years	ope of brain cancer
10. KN	contrast study	dispense medication (3 drugs/day)	3 years	vibration disamenity

Figure 3. Information of subjects. Patients in 8 and 9 showed only poor salivation. Others are Sjögren's syndrome patients.

3. Results

3.1. Variation per day of the effective salivation on the continuous vibrotactile stimulation

We examined whether or not the effective salivation occurred continuously when the vibrotactile stimulation was carried out every day. Normal subjects (26, males 15 and females 11, average age 25) used this apparatus continuously 4 or 5 days *at the same time and place* [8]. Since patients with *decreased* salivation (hyposalivation) had the psychiatric disorder in daily life, we conducted the experiment to realistically approximate the natural condition. In particular, we produced the analysis following the 89 Hz vibrotactile stimulation with 1.9 μm amplitude from the previous experiment, because this frequency produced the most effective salivation. *No gland (right and left parotid glands, and right submandibular and sublingual glands) showed a decreasing tendency with use day after day* [8].

3.2. Facial skin temperature and heart rate in normal subjects

As shown in **Figure 4A**, facial temperature under vibration apparatus was increased about 0.5°C in 2 or 3 min, and then after 15 min of *continuous stimulation*, it was up by about 0.5°C, too. *Namely, a rise in facial temperature and an increase in RR intervals (decreasing heart rate) by vibration were affected by period of stimulating time. On the other hand, by raising about 200 ms of RR intervals after 15 min an increase in heart rate was observed when the vibration was over. The reason was thought to be parasympathetic activation recoil by vibration stimulation.*

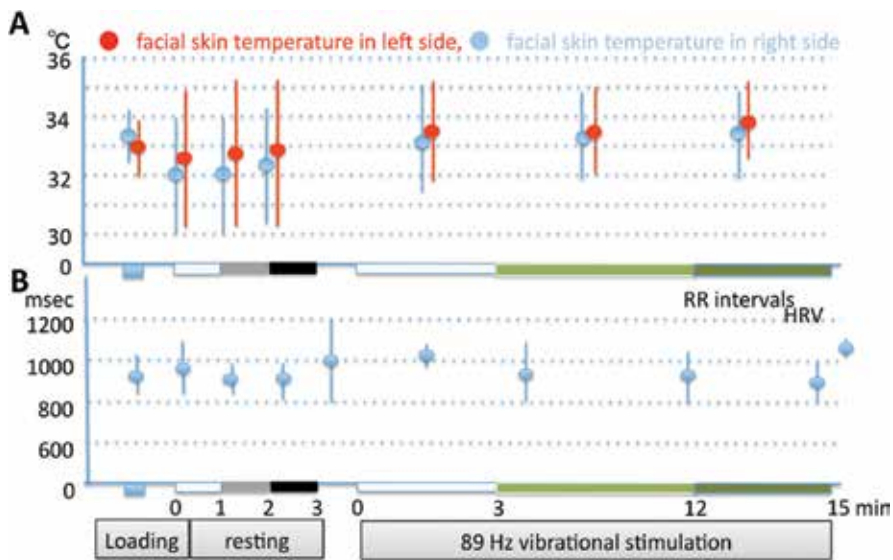


Figure 4. Facial skin temperature (A) and heart rate (B) in normal subjects.

3.3. The relationships between affected and unaffected patients after the vibrotactile stimulation

In Figure 5A, we showed changes to saliva production (Figure 5A-a) and facial temperature (Figure 5A-b) in the patients who were affected followed the vibrotactile stimulation. On the other hand, in Figure 5B, we showed changes to saliva production (Figure 5B-a) and facial temperature (Figure 5B-b) in patients who were not affected followed the vibrotactile stimulation. As shown in Figure 5A, the saliva production (elevated state from avg. 0.2 to 1.5 ml) was exponentially increased after about 5 years. The finding was shown as changes to increased face temperature (in a positive direction). However, as shown in Figure 5B, the saliva production and facial temperature remained static.

3.4. The affected and unaffected patients on the vibrotactile stimulation

In Figures 5 and 6, affected patients were divided into apocatastasis for a long period and for a short period. Affected patients with a long period were exponentially increased after about 6 (A), 5 (B) and 1.5 (D) years. Ones with a short period were exponentially increased after about 2 months (E). Facial temperature was increased with increased saliva production. On the other hand, in Figure 7, in unaffected patients, the saliva production and facial temperature remained static.

3.5. Questionnaire data

In Figure 8, we showed a survey and patients with satisfaction (good) or non-satisfaction (no/yes or bad) were examined. Patients with satisfaction had many good tendencies, but ones with non-satisfaction had many no/yes and bad.

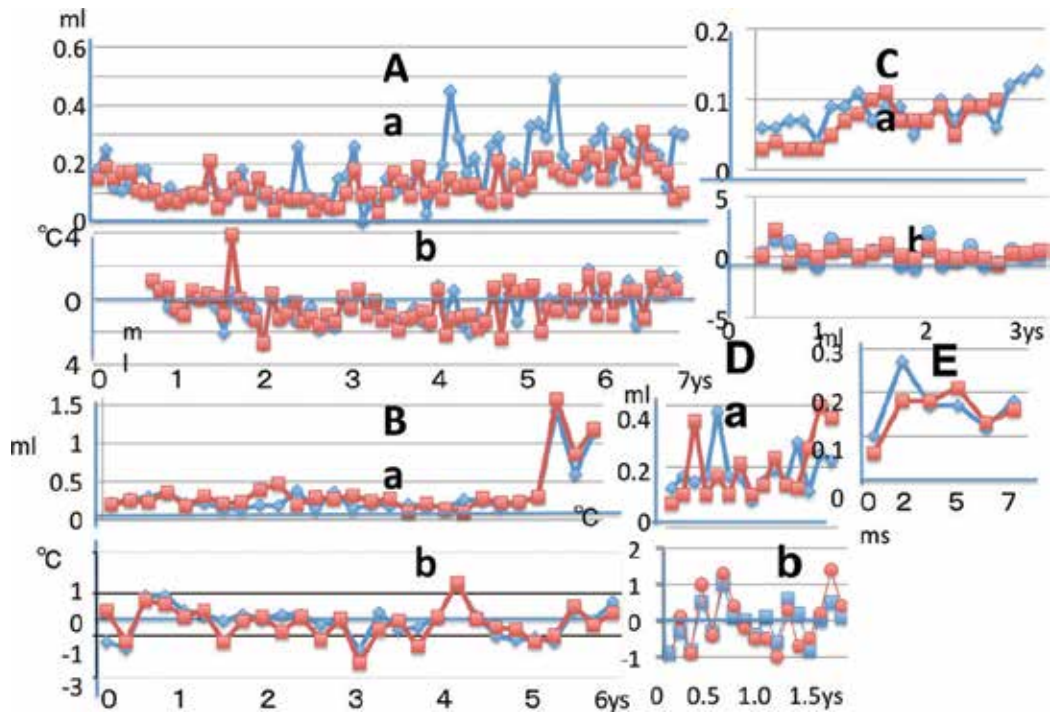


Figure 5. Patients affected by a facial vibration. A: salivation gradually increased, and temperature changed to plus tendency near 7 years. On the other hand, patients in B rapidly increased, and temperature changed to plus tendency near 6 years. Patients in C, D and E gradually increased.

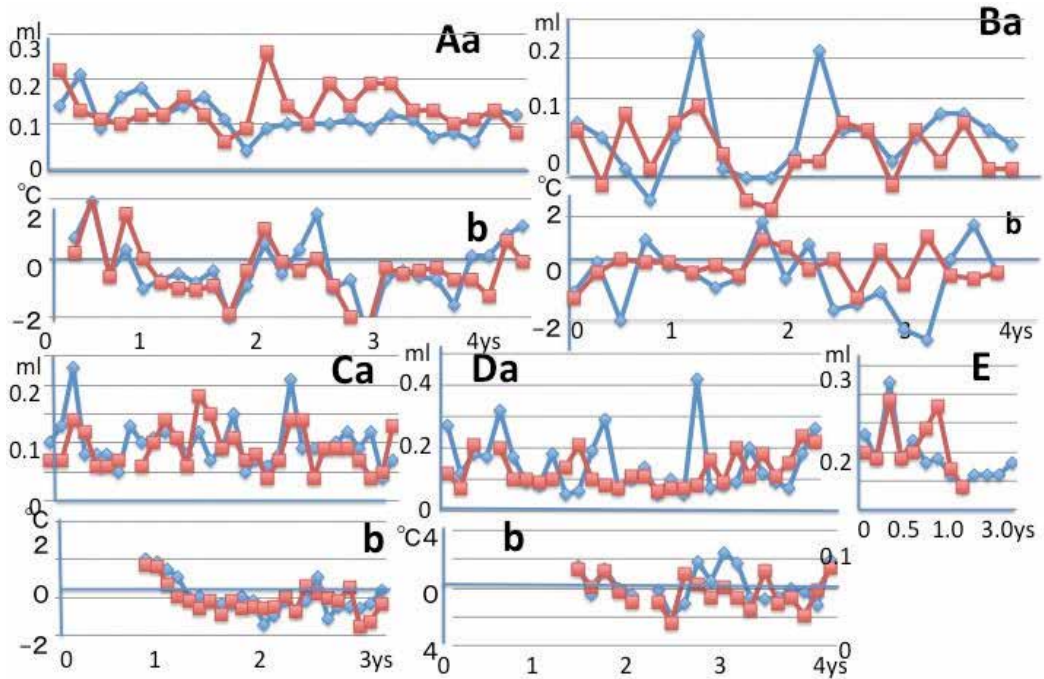


Figure 6. Patients unaffected by a facial vibration.

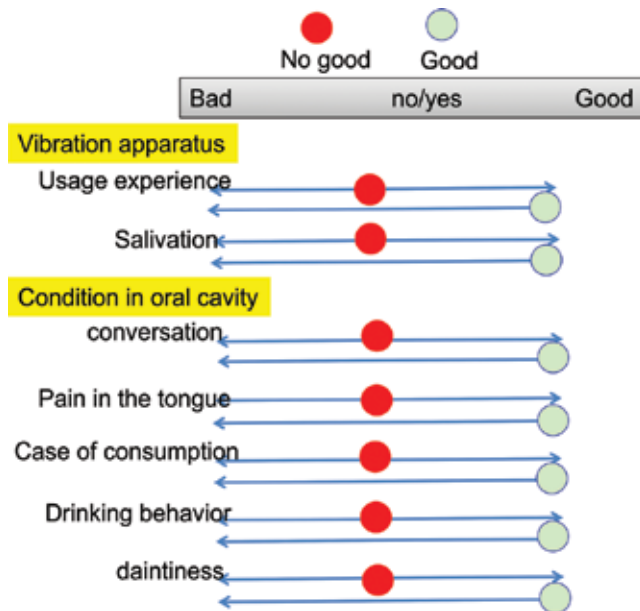


Figure 7. Typical examples of questionnaire data. We divided into three types (bad, no/yes and good) during effects of vibration apparatus and condition in oral cavity after vibration. Effect example is good (blue circles), and no-effect one is no/yes (red circles).

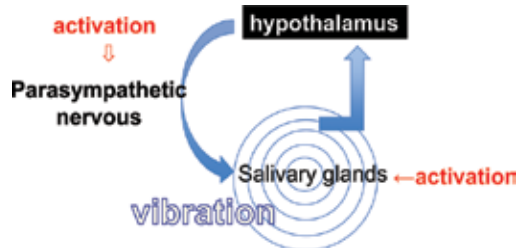


Figure 8. Summary of vibration effects. Facial vibration produces revitalization of cells of salivary glands. Furthermore, stimulation of vibration is activated at hypothalamus, and parasympathetic nerve is activated again. In particular, double activation shows 89 Hz frequency and 1.9 μ m amplitude is the most effective vibration.

4. Discussion

The continuous use of various sensory stimuli has been known to induce an adaptation [3, 7]. So we examined *the effects of vibrotactile stimulation and the adaptation in normal subjects, when the patients continuously used this apparatus every day. The patients' first desire is not to have an increase in salivary secretion from all salivary glands. However, the increase of total secretion quantities is necessary.* So, we first investigated changes in the total secretion quantities of normal subjects with the vibrotactile stimulation using a cotton roll indwelling each duct of each gland.

On parotid glands, 89 Hz vibrotactile stimuli were shown *to result in the more effective salivation in the right and left parotid glands and submandibular and sublingual glands,*

as shown in **Figure 2**. On the other hand, on the submandibular glands, 89 Hz vibrotactile stimuli with one motor were shown to result in the more effective salivation in all the glands. The findings suggest the 89 Hz vibrotactile stimuli with one motor may be the most effective salivation, and glands stimulated by vibrotactile stimuli have the tendency for the most effective salivation. Namely, on the parotid gland and on the submandibular gland vibrotactile stimuli shows the submandibular and sublingual gland, the place of stimulating portions may be the body of the mandible.

Burdette and Gale studied the effects of treatments in myofascial pain dysfunction patients [5]. Furthermore, Vrijama and Vanharantra [6] reported that discographically painful discs always produced painless feeling in the vibration examination. These facts assume that peripheral stimuli provided by vibration arrive at the central nerve (in the spinal cord and brain stem) and that these effects were exercised by the depressant effect in the cerebral cortex depending on the somatosensory information. Namely, we think that the vibration stimuli may promote the parasympathetic effects by the inhibition of sympathetic effects elicited by the pain, and so on. On the other hand, we know that the production of salivation only induces the parasympathetic effects. Furthermore, the production of salivation will be at a specific frequency and amplitude. This phenomenon may be directly produced by the vibrotactile stimulation of 89 Hz with one motor on the parotid and submandibular glands.

On the other hand, we examined the physiological characteristics of the adaptation to the vibrotactile stimulation, whether it caused decreased salivation or not. A continuous examination was performed for 4 or 5 days on 26 normal subjects [8]. Since patients with the decreased salivation (hyposalivation) are not exclusively happy in every day of their daily life, we conducted the experiment to realistically approximate the natural conditions. We did not show decreased adaptation depending on the continuous using of this apparatus, as shown in **Figure 4**. The result suggests that 89 Hz vibrotactile stimulation of the facial skin on the masseter belly may be appropriate for patients with the decreased salivation. Furthermore, we imagined mechanism of salivary production following facial vibration in **Figure 8**. Facial vibration directly activates the poor salivation of glands and then it indirectly parasympathetic nerve via hypothalamus.

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Appendices and nomenclatures

Obey neuroscience society.

Author details

Koichiro Ueda¹, Kanako Gora¹, Masaru Yamaoka², Takako Sato³, Kimiko Abe¹,
Enri Nakayama¹, Mituyasu Sato¹, Yumiko Tunoda⁴, Sumiko Akatuka⁴, Sayaka Fukui⁴,
Akinari Hayashi¹, Teruyuki Hirai¹, Sayako Ohnishi¹, Mayumi Hayata¹ and Hisao Hiraba^{1*}

*Address all correspondence to: hiraba.hisao@nihon-u.ac.jp

1 Department of Dysphasia Rehabilitation, Nihon University, School of Dentistry, Tokyo, Japan

2 Department of Basic Science, Nihon University, School of Dentistry, Tokyo, Japan

3 Department of Oral Surgery, Nihon University, School of Dentistry, Tokyo, Japan

4 Department of Dental Hospital at Surugadai, Nihon University, School of Dentistry,
Tokyo, Japan

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Combined Approaches in Sialolithiasis of Major Salivary Glands

Iordanis Konstantinidis,
Angelos Chatziavramidis and Ioannis Iakovou

Additional information is available at the end of the chapter

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Abstract

Combined (endoscopic-transcutaneous/intraoral) techniques are an effective treatment for large and/or impacted stones of the major salivary glands. This approach results in high rates of symptom improvement and gland preservation. The complication rates are relatively low, further supporting the use of these techniques as an additional tool between the classic sialendoscopy and the external classic procedures of gland removal. In this chapter, we describe the combined approach for the parotid gland and the submandibular gland and finally, the retrograde sialendoscopy through the surgical field of an open approach.

Keywords: sialendoscopy, transcutaneous, sialolithiasis, parotid, submandibular

1. Introduction

Endoscopic techniques in the management of sialolithiasis were introduced since the 1990s and gradually became the standard treatment option, decreasing the external procedures [1]. In our days only 20–25% of all symptomatic cases require an open surgical approach of the gland [1, 2]. The introduction of interventional sialendoscopy with intraductal laser fragmentation did not solve all the problems as some stones were too large to be fragmented, and some others had associated with ductal stenosis which could not be dilated [3].

In order to avoid gland removal with its associated significant morbidity, surgeons developed combined techniques (endoscopic and transcutaneous) as a solution within the frame of a gland-preserving strategy [4–6].

Indications for this combined management are failure of interventional sialendoscopy to treat impacted calculi, stones larger than 8 mm, stones located behind a stenosis which cannot be dilated, and finally a non-successful extra- or intracorporeal lithotripsy [4–8].

In this chapter, we present approaches that we use in parotid and submandibular sialolithiasis cases in order to avoid gland removal.

2. Combined approach in parotid sialolithiasis

2.1. Surgical technique

This is a procedure performed under general anesthesia. Facial nerve monitoring is mandatory as in the majority of the cases a branch of the nerve is located closely to the main ductal system.

Before the procedure, a diagnostic sialendoscopy is carried out to ensure that endoscopic localization of the stone is possible. The sialendoscope used in surgery has usually a diameter of 1.1 mm. The first step of the procedure is the identification by means of the endoscope of the stone's location in the ductal system. Then, the skin above the stone is marked as the light of the endoscope's tip can be easily detected (**Figure 1**).

The location of the stone (proximal-distal, superficial-deep in the gland) is the factor which affects the surgeon's decision regarding the incision which is required. Three incisions have been described [8]:

1. Lazy S
2. Mini parotidectomy incision extended if required to face lift
3. Straight, small incision above the stone



Figure 1. Transillumination of the parotid area and skin marking before the operation.

In the vast majority of cases, the first two incisions are used with the last one only in very superficial and/or proximal stones especially in aged people where the skin lines can hide a relatively small facial scar.

In cases where a parotid incision approach is required, the skin flap is elevated exposing enough surface of the gland for the stone removal. Transillumination of the sialendoscope's tip into the ductal system helps the precise gland-preserving dissection technique. In superficial stones, a longitudinal incision of the gland parenchyma (1–1.5 cm in length) above the endoscope light is performed (**Figure 2A**). In stones with deeper location (5 mm from the surface of the gland), a mini-flap of the gland parenchyma is prepared above the area of the ductal system which is lighted (**Figure 2B**) [9].

Stenting the ductal system is not obligatory. We suggest the use of a stent in cases where an opening of the ductal system larger than 1 cm has been performed and/or other traumatic manipulations in the duct have been done during removal of a polyp or an impacted stone. Usually, when a stent is used, this is placed in a retrograde fashion via the surgical field and secured in buccal mucosa with absorbable sutures (**Figure 3**). The stent can be left for a period of 3–6 weeks. The closure of the parotid duct is carried out using absorbable sutures. Fibrin glue and absorbable sutures are used for the repositioning of the gland flap in cases of non-superficial stone location (**Figure 4A** and **B**). A small drainage usually is needed as in parotid tumor surgery which is removed on the first postoperative day. Tight bandage for 4–5 days is proposed to avoid leak of saliva. Patients take broad-spectrum antibiotic treatment for a week and analgesics if required.

2.2. Follow-up assessment

During the first postoperative period (2–3 weeks), patients are advised to avoid nutrition which produce excessive saliva (e.g., lemon juice, etc.).

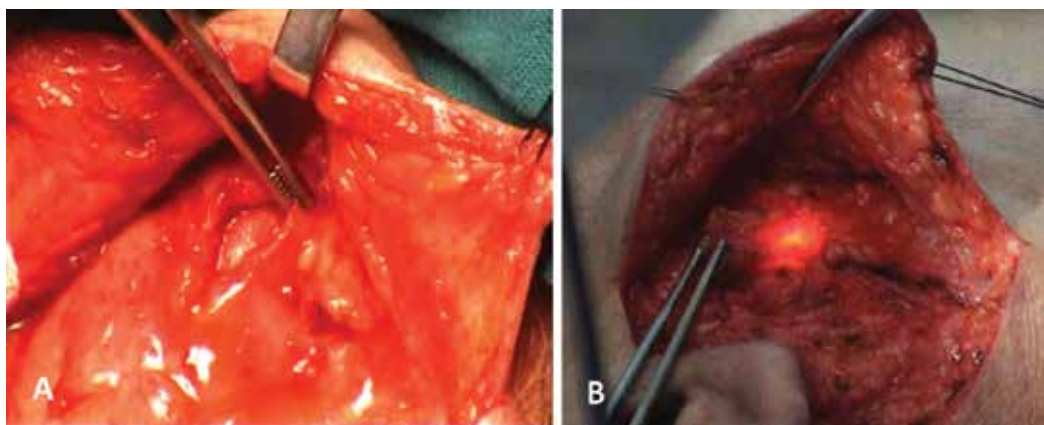


Figure 2. (A) Longitudinal incision of the gland in a superficial stone and (B) n-shaped mini-flap in a deep-located stone.

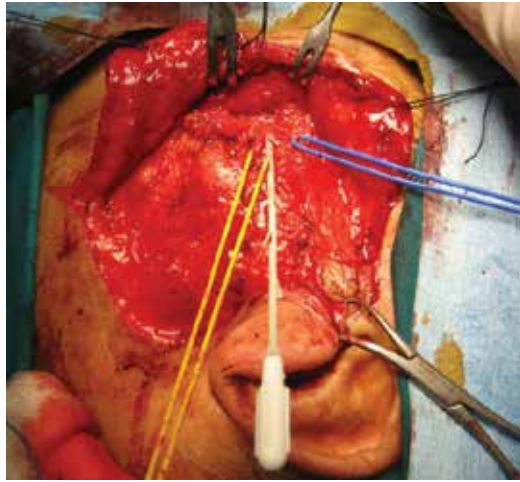


Figure 3. Stent placement in a retrograde fashion via the surgical field. The left loop isolates the main duct, and the right the facial nerve branch which run close to the ductal system.

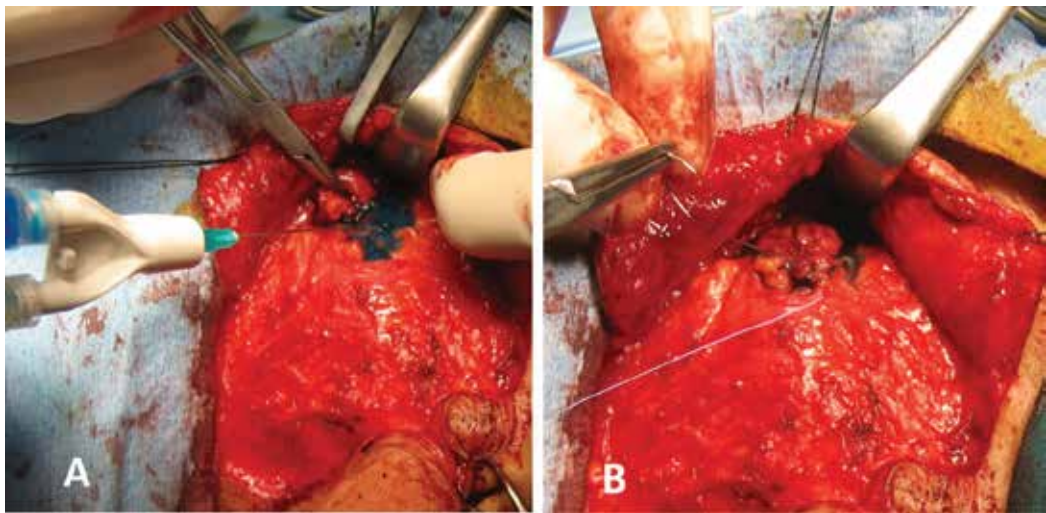


Figure 4. (A) Fibrin glue and (B) absorbable sutures are used for the repositioning of the gland flap in cases of non-superficial stone location.

Early complications such as hematoma, sialadenitis, wound infection, fistula, and obstructed and/or extruded stent have been rarely published and usually occur within the first postoperative month. In such cases, conservative management and removal of stent if needed are enough measures [9–11].

Initially, one of the surgeons' concerns about the procedure was the potential postoperative stenosis at the site of the ductal incision. This fact is expected and was confirmed in our data, as 7 patients out of 12 had an endoscopically diagnosed postoperative stenosis [9]. However,

these stenotic areas are considered without clinical impact as none of the patients suffered from postoperative swellings. The incision is parallel to the duct axon, and this may decrease the possibility for severe stenosis even in cases with extended opening. In addition stenting with precise intraoperative placement can be helpful to avoid postoperative stenosis. Undoubtedly, further studies are needed to justify the size of the ductal opening which is critical regarding postoperative stenosis.

In the same study, scintigraphic evaluation of the operated parotid glands in two phases (baseline and after stimulation) provided an objective functional evaluation 1 year after the procedures [9]. Specifically, a dynamic imaging of the whole anterior head started after a bolus intravenous injection of ^{99m}Tc . Fifteen minutes after the initial injection, diluted lemon juice was given per oz. The parameters measured were (1) uptake rate, taken as the value of the initial slope of the time-activity curve and (2) washout fraction, as a relative mobilized radioactivity from each parotid gland after the sialogogue's application. In the vast majority of cases (11/12), the procedure preserved the function of the gland with only one parotid hypofunction in a patient with long-term history of sialolithiasis.

Extracorporeal and laser intraductal lithotripsy requires expensive devices which are not always available; they are time-consuming and always have a potential risk of leaving residual stone fragments which can be a nidus for new stone formation. Moreover, some patients prefer the described surgical option as a "one shoot" intervention instead of extracorporeal lithotripsy which may need multiple sessions [10–12].

Contraindication for this procedure is the presence of diffuse ductal stenosis or multiple parenchymal stones [8, 9].

3. Combined approach in submandibular sialolithiasis

A combined approach can be also used for submandibular gland with large and/or impacted stones in a similar manner with the parotid gland.

3.1. Surgical technique

The procedure starts again with a sialendoscopy, and when the endoscope approximates, the stone then is fixed to the floor of the mouth or can be held steadily by an assistant.

Infiltration of the transilluminated area with xylocaine 2%-adrenalin 1% solution follows. The next step is an incision of the oral mucosa approximately 2 cm in length following the axis of Wharton's duct size just above the lighted area (**Figure 5A** and **B**). Caution should be taken at this point in order to avoid lingual nerve damage as it crosses Wharton's duct and this is the reason why some authors identify and isolate the nerve with a loop from the surgical field [8].

A useful surgical tip is the fact that the floor of the mouth can be pushed upward by external pressure at submandibular triangle below the patient's mandible, giving better access to the hilum area endorally.

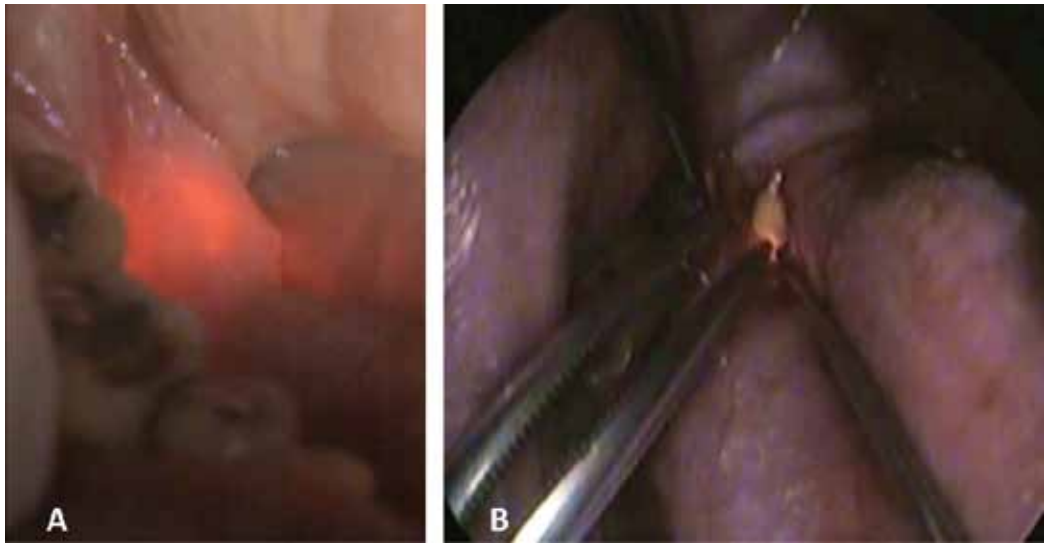


Figure 5. (A) Transillumination of the hilum area in a right submandibular gland intaraorally. (B) Incision and stone removal.

After the identification of the duct, silastic loops are positioned around the duct helping its traction and dissection. The stone is palpated by the finger or surgical instrument, and a straight incision above the stone at the axis of the duct is performed. Once the stone has been removed, an endoscopy of the ductal system behind the stone is mandatory for residual stones or strictures, etc.

The use of a stent at the end of the procedure is not universally accepted; however, it is the preference of the authors as it offers a stable floor for the ductoplasty. Its insertion through the papilla can be performed directly or with the use of the endoscope or a guide-wire for more safety especially when a precise placement to the posterior portion of the duct is required. The stent is then sutured and secured with a nonabsorbable suture to Wharton's papilla area. The time of stenting varies in the literature with a period of 3 weeks being the minimum and 6 weeks usually the maximum depending on the patients' tolerance.

3.2. Follow-up assessment

Patients receive the same instructions as in parotid stones for their diet postoperatively. Similarly, they take broad-spectrum antibiotic treatment for a week and analgesics if required with the addition of oral antiseptic solution local application after meals.

Early complications are rare and include lingual nerve damage, hematoma, and gland swelling extrusion of the stent which are usually managed in a conservative way.

A certain degree of postoperative stenosis is expected in the long term; however, the region of the hilum is large enough, and these strictures run usually as asymptomatic.

This procedure is not popular in the literature as the parotid one, because many surgeons prefer the endoral marsipulization of the duct to have access in stones at the region of the

hilum. This procedure offers equally good results although does not preserve the integrity of the ductal system. Moreover, when marsipulization of the main duct reaches the hilum, the endoscopic assessment of the residual ductal system becomes problematic due to leakage during saline irrigation.

4. Retrograde sialendoscopy

In cases with parenchymal and especially multiple stones of the submandibular gland, an external approach with removal of the gland is indicated. However, some stones may slip into Wharton's duct during surgical manipulations causing symptoms at a later time. In a study by Milton et al., authors found that 5% of patients who underwent submandibular gland removal had residual stones in the remaining duct, requiring further surgery [13].

For such cases a retrograde sialendoscopy is proposed through the surgical field after the removal of the gland [14]. A standard procedure of submandibular gland resection is performed with a transcervical incision. Identification and preservation of the lingual nerve from the gland at the area of submandibular ganglion follow along with careful dissection and skeletonization of the submandibular duct. Two stay sutures are placed on the opposing sides of the duct, proximal to the gland. After the removal of the gland, these sutures are used for stretching of the duct in order to facilitate placement of a large in outer diameter (1.4 or 1.6 mm) sialendoscope for retrograde inspection (**Figure 6**).

In cases where residual stones or debris are identified, then they can be removed by wire basket or alternatively can be pushed by the endoscope into the oral cavity (retrograde) or into the neck with the endoscope coming from the oral cavity. At the end of the procedure, stay sutures help for duct traction into the neck and its ligation proximally in order to minimize the length of the remained duct.

Contraindication of this procedure is the diffuse stenosis of the duct or severe strictures due previous trauma or surgery.



Figure 6. Retrograde sialendoscopy: insertion of the sialendoscope into Wharton's duct after removal of the submandibular gland. The duct is stretched with a silk suture.

Author details

Jordanis Konstantinidis^{1*}, Angelos Chatziavramidis¹ and Ioannis Iakovou²

*Address all correspondence to: jordan_orl@hotmail.com

1 2nd Academic ORL Department, Aristotle University, Papageorgiou Hospital, Thessaloniki, Greece

2 3rd Nuclear Medicine Department, Aristotle University, Papageorgiou Hospital, Thessaloniki, Greece

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Saliva is a complex fluid that maintains oral health and has many physiological functions. It is a noninvasive diagnostic fluid as well. Lately, salivary diagnostics has proven its potential to reach clinical practice in the near future for the early detection, diagnosis, and monitoring of various diseases. *Salivary Glands - New Approaches in Diagnostics and Treatment* is a comprehensive reference, which brings together information on salivary secretion and its disorders, the novel salivary diagnostic methods for numerous diseases, and new techniques in the treatment of salivary diseases. This book contains information for a diverse audience, including dentists, oral biologists, experimental biologists, molecular biologists, oncologists, radiologists, oral and maxillofacial surgeons, and otorhinolaryngologists.

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