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Application of a Novel Tongue Resin Replication System for *in vitro* Biofilm Studies

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Authors' contributions

This work was carried out in collaboration between all authors. Author SB designed and developed the impressed tongue resin device, performed the laboratory microorganism culture experiments, and also contributed to the writing and editing of the manuscript. Author MG designed the experiments, performed statistical analysis of the experimental dataset, and also contributed to the writing and editing of the manuscript. Author JT performed the microorganism culture count experiments and author KL designed the experiments and contributed to the writing and editing of the manuscript. All authors read and approved the final manuscript.

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ABSTRACT

Aims: Biofilms present a strategy for microorganisms to survive and resist adverse conditions. In order to further our understanding of this resistance mechanism, it is of much importance to develop new methodologies in order to assess such biofilm activities. One of the most defined biofilms is dental plaque. However, the oral cavity contains a different class of biofilm, the precise structural nature of which is dependent on its site of development. Indeed, tongue-coated biofilms have remained a focus for *in vitro* studies, especially since they represent the major source of oral malodor. The aim of this research was to investigate the growth of a biofilm on a tongue replication constructed from a resin material, and its use in further experimental investigations.

Methods: The model used for this study involved an adapted CDC biofilm reactor in order to achieve this; *Staphylococcus aureus* was used as the test organism.

Results: Biofilm culturing techniques employed demonstrated a positive growth from the microbes

retrieved from the tongue-impressed resin material. Moreover, results acquired confirmed that the resin-based tongue replication harbored significantly greater levels of *S. aureus* than those of unimpressed resin, plastic and stainless steel controls.

Conclusions: Therefore, this study provides evidence that the employment of a CDC biofilm reactor in combination with a resin tongue device creates an oral biofilm model which provides valuable information regarding our understanding of how the roughness of the dorsal lingual surface affects microbial populations in patients suffering from oral malodor.

Keywords: Biofilm; tongue coating; CDC biofilm reactor; impression material; halitosis.

1. INTRODUCTION

A biofilm is represented by a collection of microbial communities enclosed by a matrix of extracellular polymeric substances (EPSs), and separated by a network of open water channels [1]. The microbes of a biofilm are protected against environmental factors such as UV light and dehydration, in addition to host immune cells such as neutrophils and other phagocytes, together with microbicidal molecules, particularly lactams [2]. In particular, resistance to polymorphic neutrophils has been found in quorum-sensing induced P. aeruginosa, where the production of rhamnolipid B, a PMN toxin, was shown to effectively kill the phagocytic cells. Moreover, biofilm-incorporated bacteria exhibit a different phenotype from suspended bacterial cells of the same genotype [3]. Biofilms can develop on both abiotic and biotic surfaces, and act as a source of various infections, such as infective endocarditis and cystic fibrosis pneumonia [4,5].

The area of biofilm-mediated infections can include temporary or permanent medical devices such as endotracheal tubes, intravascular and urinary catheters, orthopedic implants and arterial stents, in addition to wounds, blood, and mucosal surfaces (sinuses, respiratory and genito-urinary tracts) [6–8].

One of the most studied types of biofilms are those formed within the oral cavity, in view of their interactions with biotic and abiotic surfaces that may constitute a form of chronic infection. The most common and documented oral biofilm dental plaque, responsible for dental caries (aerobic microorganisms such Streptococcus mutans). and periodontal diseases (anaerobic microorganisms such as Prophyromonas ginggivalis, Actinobacillus actinomycetiemcomitans and Treponema denticola).

A further oral biofilm that is still not well understood is that which coats on the tongue,

and which is associated with oral malodour. Indeed, firstly Yaegaki et al. [8] and subsequently Loeshe and Kazor [9], found that the dorsum of the tongue serves as the primary source of volatile sulphur compounds (VSCs). Furthermore, numerous studies have revealed positive correlations between the tongue coating status (presence and/or amount) and a range of differing parameters directly related to oral malodor, and a significant positive correlation was found between oral cavity air levels of VSCs, organoleptic scores of full mouth odor, and the presence or amount of tongue coating observed [9,10]. Moreover, de Boever [10] reported a linear Pearson correlation coefficient of 0.63 between total mouth odor and tongue coating level. Furthermore, patients presenting with a significant tongue-coating harbored bacteria than those without such tongue-coatings [11.9 x 10⁸ vs. 3.8 x 10⁶ Colony Forming Units (CFUs) per sample) [11,12].

Additionally, some researchers have described a relationship between the level of tongue-coating (semi-quantitative scale) and the degree of tongue odour assessed organoleptically [11,13]. Firstly, Quirynen et al. [11] hypnothesized that in the absence of a significant correlation between tongue coating and its microbial load, the tongue coating per se, may be responsible for oral malodor. Secondly, Mantilla et al. [13] found that the extent of a white coating on the dorsum of the tongue in periodontitis patients is greater than that observed in healthy volunteers, but they failed to detect a relationship between the presence of such a coating and salivary bacterial load [13].

These possible relationships between tonguecoatings and bacterial load were supported by the detection of benzoyl-arginine naphthylamidehydrolysing bacteria on the tongue in halitosis patients [11]. However, a further report found no association between these coatings and salivary bacterial load in healthy, gingivitis and periodontitis patients (analyzed by means of phase-contrast microscopy) [13]. However, the difficulties of studying morphological characteristics of tongue-coating biofilms are strikingly evident in the sampling process and the reproducibility of datasets acquired *in vitro*.

The aim of this research investigation was to assess the growth of a biofilm on a tongue coating replication synthesized in resin, together with its potential employment in further experimental investigations focused on this research area.

2. METHODOLOGY

In order to reproduce an in vitro biofilm, a CDC Reactor (CBR 90, Biofilm Biosurface Technologies Corporation) was utilized. This methodology was based on a continuous flow model in which growth media flowed at a specified rate, and was baffled stirred in order to generate shear force. Biofilm accumulation was then quantified by harvesting the biofilm from coupons of a known surface area, disaggregating the cells and polymeric matrix, and subsequently performing viable plate counts [14-15]. In this experimental design, this procedure was repeated with the coupon system replaced with a tongue model previously cast in resin from a tongue impression taken from a patient suffering from halitosis (Fig. 1). The patient providing the tongue impression had been enrolled voluntarily in the context of a halitosis study currently ongoing at the dental clinic in the University of L'Aquila, and approved by the University of L'Aquila Research Ethics Committee. Six replicate measurements (n = 6) were made for all materials tested.



Fig. 1. Representation of resin tongue impression

2.1 Microorganism and Culture Methods

All media was obtained from Oxoid Ltd. (Basingstoke, Hampshire, UK) unless otherwise stated. The test organism was *Staphylococcus aureus* (ATCC 6538). Strains were stored on beads (microbank PL.160/M, Prolab Diagnostics, Cheshire, UK) at -80°C. *S. aureus* was grown on Nutrient Agar (NA), and one colony per 10.0 ml volume of Nutrient Broth (NB) was grown aerobically for 24 hr. at 37°C; this culture was used to inoculate the biofilm reactor. Initial inoculi were computed via spiral plating onto NA.

2.2 Biofilm Formation

2.2.1 Batch phase

The test materials, i.e. the resin tongue, together appropriate controls (specifically the unimpressed resin tongue material, together with both plastic and stainless steel coupons) were placed in the biofilm reactor containing 500.0 ml and 1.00 ml volumes of NB and an overnight culture of S. aureus respectively. The reactor was then placed on a stirring plate (125 rpm) and incubated at room temperature for a period of 24 hr (Fig. 2). Authors underline although the tongue is internal and there may be an argument of increased temperature incubation, the ambient temperature was chosen for these studies since elevations in temperature above this level would affect growth rate, and hence also flow-rate and the overall set-up.

2.2.2 Continuous flow

Subsequent to a 24 hr. period in the batch phase, a continuous flow of media (NB) was passed through the biofilm reactor at a flow-rate of 15.24 ml/min. for 24 hr. at room temperature. The flow-rate is defined as the CDC biofilm reactor volume (350 ml) divided by residence time of the bacteria, in order to ensure that *S. aureus* numbers remain static. The doubling time of *S. aureus* was then calculated, resulting in a required residence time of 30 min.

2.2.3 Enumeration

The tongue-impressed resin and the above control materials were removed from the biofilm reactor, washed with phosphate-buffered saline (PBS) solution in order to remove planktonic cells, submerged in PBS, and then sonicated for 10 min. at 60 Hz and 100 Watts. Subsequently, the PBS solution arising therefrom was spiral-

plated onto nutrient agar and incubated at 37°C for 24 hr.; colonies were then enumerated. As noted above, relevant control materials employed for this study were resin without a tongue impression, and also plastic and stainless steel coupons.



Fig. 2. Pictorial representation of the casted tongue evaluated in a biofilm reactor system

The probe was cleansed by rinsing for 30 s. at 20,500 rpm in sterile water, followed by a rinse at an equivalent rpm value for 15 s in 70% (v/v) ethanol, and then immersed in ethanol for 1.00 min., and finally rinsed with sterile water (again at 20,500 rpm for 30 s).

2.3 Statistical Analysis

The obtained data counts were statistically analyzed by ANOVA, followed by Tukey's Honestly Significant Difference (HSD) test for individual post-hoc comparisons of the four group mean counts. The primary ANOVA p value was computed using the Welch test statistic, which allows for any inequalities in 'within-sample' variances for each of the four groups compared. A mathematical model for this experimental design is provided in equation 1, in which Yi represents the (univariate output) log₁₀ CFU/ml values observed, µ their overall population mean value in the absence of any significant, influential sources of variation, Mi the 'between-materials' source of variation, and eithe unexplained error (residual) contribution. These analyses were performed with XLSTAT2016 software modules.

$$Y_i = \mu + M_i + e_i \tag{1}$$

3. RESULTS AND DISCUSSION

The biofilm culturing techniques employed in this investigation showed a positive growth of S. aureus biofilm on the resin tongue evaluated, with a $2.91 \log_{10}$ determination of biofilm released from this material, compared with corresponding 8.9 log_{10} , 5.74 log_{10} and 5.52 log₁₀ values the smooth unimpressed resin surface, plastic and stainless steel control coupons respectively (Fig. 3). This difference is ascribable to significant obstructions to the retrieval of the biofilm presented by the rough surface of the impressed tongue resin material when expressed relative to that of a corresponding smooth (control) one, and, for the non-resin coupons, also to the differing types of materials used as substrates. The HSD test revealed that all post-hoc comparisons were significant (i.e. $p < 10^{-9}$), with the exception of the SS Control vs. Plastic Control comparison.

Since biofilms represent the most common "lifestyle" of microbes, investigations of their mechanisms of growth on different surface environments involved in human diseases are of paramount importance and hence may enable new antimicrobial agents to be developed in order to eradicate them from pathological processes. Traditionally, oral plague has always been studied in vitro or ex-vivo on similar enamel surfaces [16]. The tongue biofilm, however, has always been characterized in vitro by molecular and culturing techniques [17,18], and the models used to explore it have never considered the irregular surface on which this biofilm has the capacity to grow and proliferate (this may be ascribable to the difficulties involved in sampling processes). Importantly, the proposed model demonstrated here can be employed in ex-vivo studies, which successfully reproduce an accurate template for the oral environment.

In addition to infectious diseases, interests in biofilm formation has grown within the food industry, in particular approaches for the assessment of biofilm formation in food and water contaminants, together with metal surface corrosion and clogging. Biofilm formation has been demonstrated for numerous pathogens, and is clearly one of the main strategies for bacterial survival in a variety of sites within the human body, and also on inanimate surfaces [1]. In order to enhance our knowledge and to understand biofilm physiology, biofilm model systems are required to be developed in order to

assess these complex communities under controlled conditions [19–21].

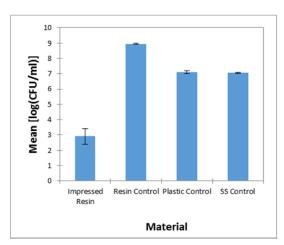


Fig. 3. Bar diagram showing plots of mean released *S. aureus* counts [log₁₀(CFU/ml)] with associated 95% confidence intervals for the four material groups tested

Abbreviations: SS, Stainless Steel

In view of the multiplicity/heterogeneity of oral biofilm populations, simultaneous multispecies evaluations of the oral microcosm are of a high level of complexity [22,23]. However, the utilisation of an *in vitro* model system with a highly complex bacterial diversity, and which also supports the growth of uncultivated oral species, is highly desirable, since it may be manipulated and explored within a controlled environment [23].

In the CDC biofilm reactor, the magnetic stirrer within the centre of the vessel provides the shear force and continuous mixing of the device's bulk fluid (introduced in the reactor by means of a peristaltic pump) over the colonised surfaces [24]. Indeed, statistical analysis performed by Goeres [25] demonstrated that the CDC biofilm reactor system serves as a reliable experimental tool to study biofilm formation evolving from a wide range of organisms.

In the reactor system, 24 identical biofilms may be formed simultaneously, and in view of the nature of the reactor components, it permits the facile removal of discs during the experimental procedure, an advantage which customizes the system to the time-dependent study of biofilm formation [26,27]. The CDC biofilm reactor model used in the study conducted by Rudney et al. [21] has been shown to be able to produce reproducible microcosm biofilms that were

representative of the oral microbiota. Notably, they were able to produce biofilm growth on disks of hydroxyapatite (HA), silorane-based substrates (LS), together with methacrylatebased composites (Z100), which were 12 mm in diameter. Basal mucin medium (BMM) was used as the growth medium, whilst saliva and dental plaque samples were utilised as inoculi. Both of the two substrates showed a positive biofilm growth, with differing microbiota compositions, and which was dependent on the differing natures of the two inoculum sources; salivary flora are more similar to that of the tongue than they are to that of dental plaque [22]. In the model used in this study, the system was adapted to accommodate a resin tongue replacing the commonly-employed surface coupon discs (Fig. 2), so that instead of 24 biofilms being produced, it was only possible to form a single one, albeit one with a larger surface

The tongue, with its large surface area, represents a unique ecological niche within the oral cavity. This promotes the retention of greater amounts of microorganisms thereon, and therefore it can harbor an extensive bacterial population. Notwithstanding, relatively little is known about the organization of the tongue flora, although it is generally accepted that they form microenvironments. The composition of the oral cavity has been shown to include Streptococci sp. and Actinomyces Porphyromonas Fusobaterium, sp., Enterobacteria [28-30].

However, correlations between the roughness of the dorsal lingual surface, tongue-coated biofilms and oral malodour biomarkers (particularly VSCs in exhaled human breath) remain to be elucidated. In view of their inner nature, the presence of fissures and grooves constitutes a reservoir for harboring microbiota, and hence provides an ecological niche for them. Interestingly, previously published research work has related the presence of deep fissures to higher total counts of bacteria (approximately a 2-fold change), and also to significantly higher mouth and tongue odour scores [10], although further investigations have failed to confirm any associations of elevated bacterial counts with an increased level of tongue surface roughness [11.13]. Similar results were obtained in this study, with microbial counts released from the control resin being much higher than that from the resin tongue impression. In order to further address this issue, many clinical trials, i.e. those involving both *in vivo* and *in vitro* models, have been postulated and explored [31-34]. However, in view of the inherent features of the tongue and biofilms thereon, the precise nature of relationships between surface roughness and the nature of such biofilms remains questionable. Notwithstanding, the results reported here demonstrate the efficacy and possible future employment of a CDC biofilm reactor, i.e. one which successfully replicates the tongue dorsal surface, for the development of a new model for assessing the oral microcosm within tongue coatings. It is therefore anticipated that this study will stimulate further investigations in this research area.

4. CONCLUSION

Although only pilot data was collected in this study, these experiments demonstrated the possibility of employing a CDC biofilm reactor in combination with a human-impressed resin tongue device in order to create an oral biofilm model. Indeed, the results acquired clearly demonstrated that the rough surface of this material presented a significant barrier to the release of *S. aureus* biofilm. This experimental approach may enhance our understanding of the mechanisms involved in the ability of the dorsal lingual surface to affect the microbial population in patients suffering from oral malodor.

ETHICAL APPROVAL

All authors hereby declare that all experiments have been examined and approved by the appropriate ethics committee and have therefore been performed in accordance with the ethical standards laid down in the 1964 Declaration of Helsinki.

COMPETING INTERESTS

Authors have declared that no competing interests exist.

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