Research Article

Study of Bacterial Dominance and its Occurrence Frequency in Dental Plaque Sample

Subramonian S¹, Segin Chandran¹, Murugan M^{2*}, Murugan T²

¹Department of Dentistry, Dr.Somervell Memorial CSI Medical College & Hospital, Karakonam - 695504, Kerala, India ²Centre for Biological Science, Noorul Islam Centre for Higher Education, Kumaracoil- 629180, Tamil Nadu, India

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ABSTRACT

The intention of the present study was to investigate the bacterial dominance of dental plaque. For this study, 20 plaque samples were collected from adult humans of the around Marthandam area, Kanyakumari District, Tamil Nadu. The collected plaque samples were inoculated separately into the Basal salt medium and Basal salt medium agar plates. The morphologically different bacterial colonies were selected, identified by studying cultural, morphological and biochemical characteristics according to Bergey's Manual of Systematic Bacteriology. As a result, 7 bacterial genera such as *Bacillus* sp. *Lactobacillus* sp. *Staphylococcus* sp-I, *Staphylococcus* sp-II, *Micrococcus* sp, *Streptococcus* sp, *Proteus* sp and *Pseudomonas* sp. were identified. Among these, *Streptococcus* sp has been found highest incidence (21.57 %) followed by *Staphylococcus* sp-I (17.65%).

Keywords: Dental plaque, Bacterial dominance, Tooth decay and Streptococcus sp

INTRODUCTION

Tooth decay (Dental caries) have plagued human since the dawn of civilization and still constitutes one of the most common human infectious disease in different parts of the world¹. It is an adherent deposit of bacteria and their products, which forms as a white greenish or even yellow film on all tooth surfaces. Dental plaque accumulates naturally at stagnant or retentive sites formed after one to two days with no oral hygiene^{2,3}. Dental caries cause destruction of enamel, dentin or cementum of teeth due to bacterial activities⁴.

Tooth decay is caused by certain types of acid producing bacteria which cause damage in the presence of fermentable carbohydrates such as sucrose, fructose and glucose⁵. Oral *Streptococci*, (*Streptococcus mutans* and *Streptococcus sobrinus*) which are the major members of oral flora, frequently cause bacteremia and infective endocarditis⁶. These microorganisms in dental plaque are embedded in a polymer matrix of host and bacterial origin. It is said to contain a diverse microbial community that may remain stable over time in a healthy state and it can harbor bacterial species that may predominate in diseased states⁷. The intention of this study was to assess the bacterial dominance of dental plaque samples.

MATERIALS AND METHODS

Collection of sample

Twenty plaque samples were collected from adult humans of around Marthandam area, Kanyakumari District, Tamil Nadu. Dental plaques from all patients were picked up through forceps and transferred into 2 ml of normal saline (sterilized). All the collected samples were stored in a cool place then transported to the laboratory.

Isolation of bacteria

The plaque samples in tubes were inoculated separately into 25 ml of Basal salt medium containing (g/L) 1.0 g yeast extract, 0.3 g (NH₄)2SO₄, 0.14 g MgSO₄·7H₂O, 0.2 g CaCl₂·2H₂O, 0.1 g NaCl, 0.05 g KH₂PO₄, 0.05 g K₂HPO₄, 0.6 mg H₃BO₃, 0.17 mg CoCl₂·6H₂O, 0.09 mg CuCl₂·2H₂O, 0.1 mg MnCl₂·4H₂O, 0.22 mg ZnCl₂, 10 g glucose⁸. The inoculated flasks were incubated at 35^oC for 48 hours. After incubation, freshly grown culture of 1 ml from each dental plaque was serially diluted up to 10⁻⁵ with distilled water. 100µl of diluted samples were spread over the Basal salt medium agar plates. The plates were incubated at 37^oC for 72 hours. After incubation, the morphologically different bacterial colonies were picked up and streaked into fresh agar plates.

Characterization and identification of bacteria

The isolated bacterial colonies were identified by studying cultural (growth on different media such as MacConkey agar, Mannital salt agar, Cetrimide agar, Eosine methylene blue), morphological (Gram's, spore and capsule staining) and biochemical characteristics (such as coagulase, catalase and oxidase test) according to Bergey's Manual of Systematic Bacteriology⁹.

RESULT AND DISCUSSION

In this present investigation, a total of 51 morphologically different bacterial isolates were obtained from plaque 20 samples. All the bacterial isolates were identified by following standard microbiological techniques such as cultural, morphological and biochemical characteristics

S	Characteria	1 0				1			
S. No	tics	S_1	S_2	S_3	S_4	S_5	S_6	S_7	S_8
110.	Gram								
1	Gram	+ve	+ve	+ve	+ve	+ve	+ve	-ve	-ve
	Morpholog								
2	Morpholog	Rod	Rod	Cocci	Cocci	Cocci	Cocci	Rod	Rod
3	y Motility	+ ve	- Ve	- Ve	- Ve	- Ve	- Ve	+ ve	+ ve
5	Cansule	i ve	ve	ve	ve	ve	ve	i ve	i ve
4	staining	- ve	- ve	- ve	- ve	- ve	+ ve	- ve	- ve
-	Spore								
5	staining	+ ve	- ve	- ve	- ve	- ve	- ve	- ve	- ve
~	Coagulase								
6	test	-	-	+ ve	- ve	- ve	- ve	-	-
	Growth on								
7	MacConke	NLF	NLF	LF	LF	LF	NLF	NLF	NLF
	y agar								
	Growth on			Golden	Golden				
8	Mannital	-	-	yellow	yellow	-	-	-	-
	salt agar			pigment	pigment				
	Growth on								
9	Cetrimide	-	-	-	-	-	-	+ve	+ve
	agar								
10	Growth on	_	_	_	_	_	_	-ve	_
10	EMB agar							ve	
11	Growth on	α-	_	ß-	ß-	ß-	α-	_	β-
	blood agar	hemolysis		hemolysis	hemolysis	hemolysis	hemolysis		hemolysis
12	Catalase	+ ve	- ve	+ ve	+ ve	+ ve	- ve	- ve	+ ve
	test								
13	Oxidase	- ve	- ve	- ve	- ve	+ ve	- ve	+ ve	+ ve
	test		~				6		5
			ts s	sn:	sn:	sp	ls s		s sl
		<u>a</u> .	llu.	000	000	swc	сп	~	na
	d v	'S S	aci	loc	loc	000	000	ls s	JMIC
	oab ttity	illu	tob_{i}	why	ikhy. I	roc	pto	teu.	лдс
	len len	ac	aci	tap v-I	tap v-L	<i>lic</i>	tre	ro	sei

Table 1: Cultural, morphological and biochemical characteristics of *Bacillus species*

'LF'= lactose fermentation; 'NLF'= Non lactose fermentation

Table 2: Percentage frequency of isolated bacterial strains

S. No.	Bacterial species	Occurrence of the Bacteria	% occurrence of the bacteria
1	<i>Bacillus</i> sp	6	11.76
2	Lactobacillus sp	4	7.84
3	Staphylococcus sp-I	9	17.65
4	Staphylococcus sp-II	5	9.80
5	Micrococcus sp	7	13.73
6	Streptococcus sp	11	21.57
7	Proteus sp	6	11.76
8	Pseudomonas sp	3	5.88
Total		51	100

(Table 1). In this study, seven bacterial genera were identified viz. *Bacillus* sp, *Lactobacillus* sp, *Staphylococcus* sp-I, *Staphylococcus* sp-II, *Micrococcus* sp, *Streptococcus* sp, *Proteus* sp and *Pseudomonas* sp. The percentage frequencies of isolated bacterial strains from 20 plaque samples were calculated. In this, *Bacillus* sp, *Lactobacillus* sp, *Staphylococcus* sp-I, *Staphylococcus* sp-II, *Micrococcus* sp, *Streptococcus* sp, *Proteus* sp and *Pseudomonas* sp were found as 11.76%, 7.84%, 17.65%, 9.80%, 13.73%, 21.57%, 11.76% and 5.88% respectively (Table 2).

In this present study, *Streptococcus* sp was found as high as higher percentage (21.57 %) of occurrence followed by *Staphylococcus* sp-I (17.65%). This result was supported

by Omolaja et al., in that study, Streptococcus mutans was found highest percentage frequency of 53.13%, followed by Staphylococcus albus (25%), Klebsiella pneumonia (9.34%). Pseudomonas sp was found at least percentage (5.88%) of occurrence⁵. This result was also supported by the results of Reyes and Dalmacio¹⁰. Nwakanma et al.¹¹ identified three bacterial strains Streptococcus, Staphylococcus and Lacto bacilli from the mouth of students. Several members of the Streptococcaceae family were identified in the saliva and plaque and they are as considered initial colonizers of the oral cavity¹². The results of this study confirmed previous reports that some *Streptococcus* species are associated with healthy states¹². The most commonly associated and strongly implicated member of this family with dental caries is Streptococcus mutans¹⁴.

CONCLUSION

It was isolated and identified, 8 genus of bacterial strain includes *Bacillus* sp, *Lactobacillus* sp, *Staphylococcus* sp-I, *Staphylococcus* sp-II, *Micrococcus* sp, *Streptococcus* sp, *Proteus* sp and *Pseudomonas* sp were highly prevalent in dental caries and among these, *Streptococcus sp* has been found highest frequency.

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