ENTEROBACTERIACEAE AND PSEUDOMONADACEAE ON THE DORSUM OF THE HUMAN TONGUE

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ABSTRACT

Objective: The aim of this study was to correlate the presence of Enterobacteriaceae, Pseudomonadaceae, Moraxellaceae and Xanthomonadaceae on the posterior dorsum of the human tongue with the presence of tongue coating, gender, age, smoking habit and denture use. Material and Methods: Bacteria were isolated from the posterior tongue dorsum of 100 individuals in MacConkey agar medium and were identified by the API 20E system (Biolab-Mérieux). Results: 43% of the individuals, presented the target microorganisms on the tongue dorsum, with greater prevalence among individuals between 40 and 50 years of age (p = 0.001) and non-smokers (p=0.0485). Conclusions: A higher prevalence of Enterobacteriaceae and Pseudomonadaceae was observed on the tongue dorsum of the individuals evaluated. There was no correlation between these species and the presence and thickness of tongue coating, gender and presence of dentures.

Key words: Enterobacteriaceae. Pseudomonadaceae. Oral cavity. Tongue.

INTRODUCTION

The microbial composition of dental biofilm has been extensively studied. However, fewer studies have evaluated the microbiota of the oral soft tissue which comprises at least 80% of the surfaces that can be colonized within the mouth¹⁰. Although the tongue dorsum appears to shelter a complex microbiological niche, knowledge of the role of tongue microbiota in systemic and oral diseases is very limited.

There has been increased interest in the study of the microbiota of tongue, due to its possible association with oral halitosis^{7,14,24}. Previous studies have suggested that Enterobacteriaceae may be involved in halitosis, as they produce putrefied gases when they metabolize proteins or certain amino acids⁵. Nasal and paranasal secretion may accumulate in the posterior region of the tongue dorsum, favoring the formation of tongue coating causing halitosis as a result of amino acid degradation by bacteria under anaerobic conditions¹⁵.

The importance of tongue as a potential microbial reservoir has been suggested by some studies. A study of the ultrastructure of the morphotypes of bacteria residing to the tongue's surface showed the presence of 35 different

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bacterial morphotyes¹. Tanner, et al.²⁵, by using checkerboard DNA probe assay, verified that some cariogenic or periodontal pathogens microorganisms were more frequently isolated from tongue than tooth samples in children under 18 months, and suggested that the tongue is an important microbial reservoir. This role was also reported among adults and previous studies revealed that tongue might have a role in the recolonization of dental surfaces and be a reservoir of periodontal pathogens³.

Microorganisms of the Enterobacteriaceae family and *Pseudomonas* genus may act as opportunist pathogens and colonize the human oral cavity, especially in patients with debilitating diseases who are submitted to prolonged treatments with antibiotics or cytotoxic medications^{17,18,20,22}. According to Zhu, et al.²⁷ the oral reservoir of coliforms in people suffering from stroke is noteworthy as *K. pneumoniae* may cause aspiration pneumonia.

Previous studies reported the presence of this group of microorganisms in the oral cavity. A clinical study conducted by Goldberg, et al.⁵ evaluated the prevalence of Enterobacteriaceae in four different populations and they detected Enterobacteriaceae in 48% of patients with complete dentures, 27.1% of patients with halitosis, 16.4% of controls and 13% of orthodontic patients. The species

Klebsiella oxytoca, Klebsiella pneumoniae pneumoniae, Enterobacter gergoviae and *Enterobacter cloacae* were isolated from the tongue.

Increased counts of coliforms were observed among individuals with chronic nail-biting habit² and after the installation of fixed orthodontic appliances⁶. On the other hand, no differences in oral carriage of coliforms were observed among stroke sufferers²⁷ and HIV-infected patients²⁶. During chemotherapy, Napeñas, et al.¹² reported that the most frequent Gram-negative species isolated were from the Enterobacteriaceae family, *Pseudomonas* spp. and *E. coli*.

Considering the importance of the oral cavity as a reservoir of potentially pathogenic microorganisms, the aim of this study was to analyze the presence of bacteria of the Enterobacteriaceae, Pseudomonadaceae, Moraxellaceae and Xanthomonadaceae families in the posterior region of the tongue dorsum in a population of adult individuals. Also, the study aimed to correlate of the prevalence of these microorganisms with the presence of tongue coating, gender, age, smoking habit and denture use.

MATERIAL AND METHODS

The research protocol was approved by the local Research Ethics Committee (UNITAU/Protocol 163/05) and the volunteers signed an informed consent form prior to entering the study. One hundred individuals of both genders (50 females and 50 males) between 30 and 50 years of age were selected and evaluated with regard to the presence of tongue coating and dentures. The smoking habit, period of time from last oral hygiene, tongue cleaning habit, and whether the volunteer was capable to identify him/herself as having halitosis was evaluated by means of a questionnaire. Individuals who wore orthodontic appliances and used antibiotics or oral rinses within 6 months prior to the study were excluded.

The presence of coating on the tongue dorsum was quantified accordance with Tanaka, et al.²⁴. The following scores were attributed to the tongue coating area: 0) none; 1) coating on the posterior third; 2) coating on the posterior and middle thirds; and 3) coating on the posterior, middle, and anterior thirds of the tongue dorsum. Tongue coating thickness was defined by the following scores: 0) none; 1) thin layer with visible papillae; 2) moderate layer with some visible papillae; and 3) thick layer with no visible papillae.

Tongue coating was evaluated by a single calibrated examiner, after the percentage of agreement had been assessed. Initially, the area and thickness of tongue coating in 10 volunteers were measured, and after 24 h, the same volunteers were evaluated again (k = 0.9).

Material was collected from the tongue dorsum in the region of the circumvallated papillae with a calibrated sterile disposable 10 µL loop (Inlab Diagnostic; Diadema, SP, Brazil). Sampling consisted in collecting material by scrapping the entire region until loop content was completed. Then, the material was transferred to safe-lock micro test tubes (Eppendorf, Barkhausenweg, Hamburg, Germany) containing 1 ml of sterile PBS (phosphate buffered saline; pH 7.4), and homogenized for 30 s (Tube agitator, AP 56, Phoenix, Araraquara, SP, Brazil). An aliquot of 0.1 mL of this suspension was plated on MacConkey agar media plates (Difco, Detroit, MI, USA) in duplicate. The plates were incubated at 37°C for 48 h and the number of colony forming units (cfu)/mL was calculated. Then, 5 colonies representative of the different colonial morphology observed were isolated and stored in gelose agar at room temperature until identification. The strains obtained were identified using the API 20E system (Biolab-Meriéux, France).

The chi-square test was applied to correlate the positive and negative cases with tongue coating area and thickness (α =0.05). Fisher's exact test was used to compare the positive and negative cases with gender, age group, smoking habit, and presence of dentures. Statistically significant difference was considered when p < 0.05.

RESULTS

Forty-three percent the individuals showed one or an association of the studied bacterial families (Enterobacteriaceae, Pseudomonadaceae, Moraxellaceae and Xanthomonadaceae) on the tongue dorsum. The isolated presence of Enterobacteriaceae and Pseudomonadaceae was observed among 29% and 6% of the volunteers, respectively. Higher prevalence was observed among individuals between 40 and 50 years of age (Table 1) and non-smokers (Table 2). No statistically significant (p=0.053) associations were found between the prevalence of these microorganisms and gender.

Six volunteers wore complete dentures and out of these, 4 presented Enterobacteriaceae and/or Pseudomonadaceae. Nine wore removable partial dentures, and of these 6 were

TABLE 1- Number of volunteers that presented Enterobacteriaceae, Pseudomonadaceae and Moraxellaceae on the tongue dorsum according to the age group

Age group	Positive Cases		Negative Cases		Total	
	n	%	n	%	Ν	
30-39	16	37.2	40	70.17	56	
40-50	27	62.8	17	29.83	44	

Fisher's exact test (p=0.001).

positive for the studied microorganisms. Fourteen wore fixed dentures, and of these 6 presented Enterobacteriaceae and/ or Pseudomonadaceae. No statistically significant association was found in the prevalence of the studied families and the presence of dentures (p=0.890).

Combined presence of Enterobacteriaceae and Pseudomonadaceae was observed in 6% of the sample, and the association of Enterobacteriaceae, Pseudomonadaceae and Moraxellaceae was present in 1% of the population (data not shown).

A total of 63 isolates were obtained from 43 positive individuals. *Enterobacter cloacae* was the most prevalent strain (22.3%), followed by *Pasteurella pneumotropica*/ *haemolytica* (14.3%). Of the Pseudomonadaceae family, *Chryseomonas luteola* was the most prevalent species (9.5%) (Table 3).

The simultaneous presence of two or more species on the tongue dorsum was observed in 16 volunteers. *E. cloacae* was the species that appeared more frequently in association with other species, such as *Chryseomonas luteola*, *Pantotea* spp., *Flavimonas oryzihabitans*, *Klebsiella pneumoniae*, *Yersinia enterocolitica*, *Shigella* spp. and *Pasteurella pneumotropica*. Other combinations included *Klebsiella oxytoca* and *C. luteola*, *Klebsiella oxytoca* and *Serratia plymuthica* and *Shigella* spp. and *K. pneumoniae rhinoscleromatis*.

TABLE 2- Number of volunteers that presented Enterobacteriaceae, Pseudomonadaceae and Moraxellaceae on the tongue dorsum according to the smoking habit

Smoking habit	Positive Cases		Negative Cases		Total	
	n	%	n	%	Ν	
Smoking	6	13.95	14	24.56	20	
No smoking	37	86.05	43	75.44	80	

Fisher's exact test (p=0.0485)

TABLE 3- Frequency of isolation and mean log of colony forming unit per milliliter (cfu/mL) for the Enterobacteriaceae, Pseudomonadaceae, Moraxellaceae and Xanthomonadaceae species on the tongue dorsum of the 100 volunteers

Species	Frequency	Mean log of cfu/ml	
·	n	%	
Enterobacter cloacae	14	22.3	2.50
Pasteurella pneumotropica/haemolytica	9	14.3	4.14
Pantoea spp. 1	6	9.5	3.18
Chryseomonas luteola	6	9.5	3.79
Flavimonas oryzihabitans	3	4.7	3.46
Klebsiella oxytoca	3	4.7	2.69
Klebsiella pneumoniae rhinoscleromatis	3	4.7	3.90
Shigella spp.	3	4.7	4.19
Pantoea spp. 2	2	3.2	1.54
Pseudomonas aeruginosa	2	3.2	4.93
Yersinia enterocolitica	2	3.2	4.08
Klebsiella pneumoniae pneumoniae	2	3.2	2.84
Acinetobacter baumannii/calcoaceticus*	1	1.6	1.00
Enterobacter cancerogenus	1	1.6	2.55
Klebsiella ornithinolytica	1	1.6	2.50
Klebsiella pneumoniae ozaenae	1	1.6	3.20
Pseudomonas fluorescens/putida	1	1.6	2.85
Serratia marcescens	1	1.6	1.60
Serratia plymuthica	1	1.6	1.60
Stenotrophomonas maltophilia**	1	1.6	2.14
Total	63	100	2.93

* Moraxellaceae family; **Xanthomonadaceae family.

Area	Positive	Positive Cases		Negative Cases	
	n	%	n	%	Ν
0	7	16.28	10	17.55	17
1	26	60.46	38	66.66	64
2	8	18.6	9	15.79	17
3	2	4.66	0	0	2

TABLE 4- Number of volunteers that presented Enterobacteriaceae, Pseudomonadaceae and Moraxellaceae on the tongue dorsum according to the tongue coating area

Chi-square Test (χ^2 = 2.936), (p=0.4016).

TABLE 5- Number of volunteers that presented Enterobacteriaceae, Pseudomonadaceae, Moraxellaceae and Xanthomonadaceae on the tongue dorsum according to the tongue coating thickness

Thickness	Positive	Positive Cases		Negative Cases	
	n	%	n	%	Ν
0	7	16.28	10	17.55	17
1	14	32.56	15	26.32	29
2	21	48.84	30	52.63	51
3	1	2.32	2	3.5	3

Chi-square test ($\chi^2 = 0.536$), (p=0.911).

Regarding tongue coating, a high percentage of subjects with scores 1 for coating area and scores 1 and 2 for thickness was observed (Tables 4 and 5). No significant associations among scores of coating area (p=0.402) or thickness (p=0.911) and detection of microorganisms were found.

DISCUSSION

In this study, the prevalence of Enterobacteriaceae and/ Pseudomonadaceae, Moraxellaceae or and Xanthomonadaceae in the posterior region of the tongue dorsum was 43%. Studies conducted in Brazil reported similar values, though they analyzed oral rinse samples. Santos, et al.¹⁷ observed that the prevalence of these microorganisms in the oral cavity of individuals with chronic periodontitis was of 43.18%, and in another study, Santos and Jorge¹⁶ observed higher prevalence of Enterobacteriaceae and/or Pseudomonadaceae in the oral cavity (51%). These results might indicate that, as suggested by previous studies^{3,25}, tongue can be the primary reservoir of microorganisms from these families.

Out of the positive volunteers in this study, 69.05% presented Enterobacteriaceae in posterior dorsum of the tongue. On one hand, there was a significant difference between the prevalence observed in our study and other analyzing the same age group. Sedgley and Samaranayake¹⁹, in a study conducted in Hong Kong with oral rinse samples (32%). Goldberg, et al.⁵ analyzing saliva, periodontal pocket

and tongue dorsum samples observed a prevalence of Enterobacteriaceae of 27.1% in patients with halitosis. Also, Leung, et al.⁸, evaluated mouth rinse samples of individuals after radiotherapy treatment at head and neck region, and observed that the prevalence of Enterobacteriaceae in individuals between the ages of 48 and 60 years was 32%. On the other hand, the results of the present study were similar to those obtained by Leung, et al.⁹ for individuals > 60 years of age (62.5%). More studies in the age group of >60 years in a Brazilian population may clarify these differences.

There was greater prevalence of positive cases for Enterobacteriaceae and Pseudomonadaceae in the age group from 40 to 50 years of age in the present study. Sedgley and Samaranayake¹⁸ also observed prevalence of Enterobacteriaceae with statistically significant results in the older age groups (over 50 years of age). In a study with volunteers from 43 to 60 years of age and over 60 years of age, Leung, et al.⁹ found higher prevalence of Enterobacteriaceae in the volunteers over the age of 60 years. Further research analyzing other age groups is necessary.

In this study, no statistically significant associations were found between the prevalence of these microorganisms and gender. Similar results have been reported elsewhere^{16,19}. On the other hand, higher prevalence of positive cases among males (p<0.05) was observed in another study¹⁷.

Previous studies reported the role of complete dentures as a potential reservoir of respiratory pathogens and facilitating factor for the colonization of the oropharynx^{22,23}, in this study no correlation between the presence of microorganisms and dentures could be proven. This result may be related to the small number of dentures users in the sample. Future studies including larger population of denture users could better clarify this correlation.

Enterobacter cloacae was the most frequently isolated Enterobacteriaceae in this study, corroborating the data reported by Sedgley and Samaranayake¹⁹, Santos and Jorge¹⁶, Santos, et al.¹⁷ and Leung, et al.⁸. On the other hand, Galili, et al.4, studying leukemia patients, observed that the most isolated Enterobacteriaceae was Klebsiella followed by Enterobacter. The most frequently isolated Pseudomonadaceae species was Chryseomonas luteola, which was found in 9 volunteers, with a prevalence of 9.38%. Chryseomonas luteola was found in 1 patient with a prevalence of 2.08% in a previous¹⁷. However, according to Leung, et al.9 the most prevalent Pseudomonadaceae in adult oral cavity was Pseudomonas fluorescens/putida (5.3%) and Stenotrophomonas maltophilia (5.3%).

The important correlation of *K. pneumoniae* in the oral cavity and risk of aspiration pneumonia in susceptible individuals was cited recently²⁷. In the population of stroke sufferers studied by these researchers, this species was the most frequently identified coliform bacteria. In the present study, carried out among volunteers without any systemic disease, this species was observed in small proportion of the population (3.12%).

Out of the 100 volunteers analyzed in the present study, the greatest prevalence of tongue coating occurred in the posterior third of the tongue. Similar studies^{11,13}, also observed the greatest prevalence of tongue coating in the posterior region of the tongue dorsum. The anterior two thirds of the tongue dorsum are constantly rubbed by the hard palate, possibly cleaning the tongue²¹. The posterior third of the tongue is in contact with the soft palate, where there is no roughness and cleaning effect, and thus it is more prone to accumulate tongue coating. According to Rosenberg¹⁵, nasal and paranasal secretions may accumulate in the posterior region of the tongue dorsum, favoring the formation of tongue coating.

There was no statistically significant relation between the area and thickness of tongue coating and the presence of Enterobacteriaceae and/or Pseudomonadaceae in this study. Mantilla Gomez, et al.¹¹ did not find any significant difference between the coloring and thickness of tongue coating in the dorsal region and bacterial load in saliva samples.

CONCLUSIONS

According to the data of the present study, it may be concluded that the prevalence of individuals positive for Enterobacteriaceae and/or Pseudomonadaceae in the posterior region of the tongue dorsum was high, especially among volunteers in the 40-50-year-old age group and nonsmokers. There was no significant correlation between the presence of Enterobacteriaceae and Pseudomonadaceae and tongue coating, gender or denture use.

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