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## Tongue Piercing: The Impact of Material on Microbiological Findings

 Ines Kapferer, M.D.<sup>a,\*</sup>, Ulrike S. Beier, M.D.<sup>a</sup>, Rutger G. Persson, Ph.D.<sup>b,c</sup>
<sup>a</sup> Department of Restorative and Operative Dentistry, Dental School, Innsbruck Medical University, Innsbruck, Austria

<sup>b</sup> Laboratory of Oral Microbiology, School of Dental Medicine, University of Bern, Bern, Switzerland

<sup>c</sup> Department of Periodontology, School of Dental Medicine, University of Bern, Bern, Switzerland

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### A B S T R A C T

**Purpose:** Biofilms on oral piercings may serve as a bacterial reservoir and lead to systemic bacteremia or local transmission of pathogenic microbiota. The use of piercing materials which are less susceptible to biofilm accumulation could contribute to prevention of problems. The present study investigated whether there are microbiological differences in bacterial samples collected from tongue piercings made of different materials.

**Methods:** A total of 85 subjects with tongue piercings participated in this study. After a baseline dental examination, sterile piercings of four different materials were randomly allocated to the study subjects. After 2 weeks, microbiologic samples were collected and processed by checkerboard deoxyribonucleic acid-deoxyribonucleic acid hybridization methods.

**Results:** About 28.8% of subjects reported 61 lingual recessions ( $1.91 \pm .96$  mm), whereas 5% reported tooth chipping on one tooth each. With the exception of *Aggregatibacter actinomycetemcomitans* (Y4), *Fusobacterium nucleatum* species, and *Parvimonas micra*, bacteria associated with periodontitis were not commonly found in the samples from studs or piercing channels. Of the 80 bacterial species, 67 were found at significantly higher levels ( $p < .001$ ) in samples from stainless steel than from polytetrafluoroethylene or polypropylene piercings.

**Conclusion:** The low bacterial counts from piercing channels suggest that having a tongue pierced would not contribute to an increased risk for oral infection. The present study demonstrated that studs made of steel might promote the development of a biofilm, whereas those made of polytetrafluoroethylene or polypropylene may be rather inert to bacterial colonization. The finding of *Staphylococci* on steel and titanium studs may suggest an elevated risk for complication if the piercing channel is infected.

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Body piercing and other body modifications have increased tremendously in popularity in recent years [1], especially among teenagers and young adults in the industrial world. Oral piercing mostly involves the lips, tongue, and cheeks, with tongue being the most commonly pierced intraoral site [2,3]. In a cross-sectional household survey of 10,503 British adults, the prevalence of tongue piercing was 6.5% in those aged 16–24 [3]. From a medical perspective, the use of tongue jewellery cannot be considered a harmless fashion trend as it can produce undesired local and general effects [4]. Early complications include bacterial infec-

tion, pain, swelling, prolonged bleeding, and difficulties in swallowing, speech, and mastication [5]. Late complications include chipped and fractured teeth, gingival trauma, localized periodontitis, persistent difficulties in oral functions, and swallowing of the device [5]. The published data on medical implications of tongue piercing mainly includes case reports and a limited number of clinical studies [4–7]. Therefore, many biological questions related to these foreign bodies, such as biofilm formation, remain unaddressed. With infections being one of the most frequent piercing complications [8], biofilm formation on oral piercing is a fundamental issue. Additionally, biofilms on oral piercings may serve as a bacterial reservoir and lead to systemic bacteremia and even septic complications. The piercing procedure exposes the piercee to a high risk of infection because the oral cavity harbors a huge amount of bacteria [4]. The high

\* Address correspondence to: Ines Kapferer, M.D., Department of Restorative and Operative Dentistry, Dental School, Innsbruck Medical University, Reichenauerstraße 46, 6020, Innsbruck, Austria.

E-mail address: ines.kapferer@gmx.net

vascularity of the area is a further aspect to be considered [4]. Although complications such as infective endocarditis [9], epidural abscess [10], chorioamnionitis [11], herpes simplex virus hepatitis [12], hepatitis C virus infection [13], toxic shock syndrome [14], and cerebellar brain abscess [15] are rare, they are dangerous complications. Additionally, biofilms on oral piercings may serve as reservoirs for bacteria associated with periodontitis, because of the anaerobic condition in the piercing channel [4]. Thus, the use of piercing materials less susceptible to biofilm accumulation could contribute to alleviation or even prevention of problems. Currently, there are no data on the additional role of the piercing material in plaque accumulation on oral piercings.

Stainless steel (SS), titanium (Ti), polytetrafluoroethylene (PTFE), and the polypropylene Borealis A/S Bormed HD810MO (PP) are commonly used as piercing materials. The surfaces of SS as well as Ti are well known for good mechanical properties, high corrosion resistance, and excellent biocompatibility [16]. PTFE is an autoclavable synthetic polymer consisting of carbon and fluorine [17]. Bormed is a heat and radiation sterilizable PP homopolymer designed for medical applications [18]. These four piercing materials differ in surface roughness  $R_a$  (which is highest for PTFE), in wettability (which is lowest in PTFE), and surface chemistry (unpublished observations). Materials like gold and silver – popular in other body regions – are only of scarce usage with regard to piercing in the oral cavity, and thus they have not been considered in this study.

The present study aimed to assess microbiological findings in association with tongue piercing in a population obtained from a nondental setting. It was hypothesized that there are microbiological differences in bacterial samples collected from tongue piercings made of different materials. It was also hypothesized that the piercings carry the same characteristic bacteria as found in the piercing channels and that the biofilm on the tongue is independently similar to the other study locations.

## Methods

### Ethical considerations

The Ethics Committee of Innsbruck Medical University, Austria, approved the study. The study was conducted in accordance with the Helsinki Declaration of 1975, as revised in 2000. All subjects signed informed written consent before investigation. The study was performed in 2008 at the Department of Operative and Preventive Dentistry, Innsbruck Medical University, Innsbruck, Austria. At the conclusion of the clinical examination, participants obtained appropriate compensation, and were informed about their oral status and any diagnosed muco-gingival lesions. Subjects with diagnosed pathological conditions were offered appropriate treatment.

### Study subjects

Posters and flyers were dispersed on the university campus, in high schools, and vocational schools in Innsbruck, Austria, to recruit subjects for this study. The study cohort included 85 subjects with tongue piercing. The piercing had to be in situ for at least 6 months. The following exclusion criteria were applied: pregnancy and lactating women, medication with an effect on gingival tissues, antibiotic medication in the last 6 months or need for antibiotic prophylaxis, chlorhexidine use in the last 6

months, nonplaque induced gingival disease, and earlier diagnosis of periodontitis.

### Questionnaire

Participants were asked to complete a questionnaire to determine demographic and medical data, smoking habits, characteristics of the piercing device worn, and postpiercing complications. If tooth chipping was found during clinical examination, subjects were asked to provide information about the circumstances under which the chipping occurred.

### Clinical examination

Clinical periodontal conditions were recorded at six sites per tooth, excluding wisdom teeth. Probing depth (PD) was measured with a pressure-calibrated probe (ClickProbe 1395, KerrHawe, Bioggio, CH) to the nearest millimeter. Bleeding on probing (BOP) [19] was recorded dichotomously. Presence or absence of plaque was measured using the plaque control record [20]. The amount of recession was measured from the cemento-enamel junction to the free gingival margin at six sites per tooth. Clinical attachment level was calculated by adding the amount of recession and PD. One investigator (I.K.) performed all measurements and collected all samples.

After the periodontal examination, the personal piercings of the study subjects were substituted by one of the test piercings of four commonly used piercing materials: Ti, SS, PP, PTFE. Randomization was performed before experiments by computer-generated randomization (Excel, Microsoft Corp., Redmond, WA), and piercing materials were allocated to the study subjects in the chronological order of appointment. The length of the stud for each subject was measured. Piercings were cut to the appropriate length for PP and PTFE. Different lengths of piercings were available to the investigator for Ti and SS. All piercings were packaged individually and sterilized (121°C, 20 minutes, steam autoclave Belimed KHS 2000 [Belimed Sauter AG, Sulgen, CH]) before the experiments. Packages were only opened at the visit when the devices were placed and carefully handled to prevent contamination during clinical manipulations. Special attention was paid to prevent damage to the channel tissue when removing and inserting the studs. Study subjects had the piercing 2 weeks in situ. The test piercings were then removed. The piercings with adhering biofilms were placed in 1.5 mL PBS in deoxyribonuclease free laboratory tubes (natural flat cap microcentrifuge tubes, Starlab GmbH Ahrensburg, Germany), and sonicated for 30 seconds to disperse adhering bacteria. The efficacy of the removal of the bacteria has been tested in preliminary studies through scanning electron microscopy. The studs were removed and the solutions were assayed for bacterial identification. At the laboratory, .15 mL Tris ethylenediaminetetraacetic acid (EDTA) buffer (10 mM Tris-HCl, 1.0 mM EDTA, pH 7.6) and .10 mL .5 M NaOH were added to each Eppendorf tube. A swab was used to collect microbiological samples from the tongue. An endodontic paper point size 55 (Absorbent Paper Points, Dentsply/Maillefer, Ballaigues, CH) was inserted in the piercing channel and kept in situ for 20 seconds. Efforts were made to move the paper point against the channel linings of the piercing locations after the stud had been removed. The collected paper points were placed in individual dry Eppendorf tubes (1.5-mL natural flat cap deoxyribonucleic acids (DNAs) and ribonucleic acids free micro-centrifuge tubes, Starlab, Ahrensburg, Germany).

**Table 1**  
Reference bacteria strains included in the DNA-DNA checkerboard analysis

Species	Collection	Species	Collection
<i>Actinomyces israelii</i>	ATCC 12102	<i>Lactobacillus jensenii</i>	GUH 160339
<i>Actinomyces naeslundii</i> (type I + II)	ATCC 43146	<i>Lactobacillus vaginalis</i>	GUH 078092
<i>Actinomyces neuii</i>	GUH 550898	<i>Leptotrichia buccalis</i>	ATCC14201
<i>Actinomyces odontolyticus</i>	ATCC 17929	<i>Mobiluncus curtisii</i>	GUH 070927
<i>Aggregatibacter actinomycetemcomitans</i> (a)	ATCC29523	<i>Mobiluncus mulieris</i>	GUH 070926
<i>Aggregatibacter actinomycetemcomitans</i> (Y4)	ATCC 43718	<i>Neisseria mucosa</i>	ATCC 33270
<i>Aerococcus christensenii</i>	GUH 070938	<i>Parvimonas micra</i>	ATCC 19696
<i>Aanaerococcus vaginalis</i>	GUH 290486	<i>Peptoniphilus</i> sp.	GUH 55097
<i>Atopobium parvulum</i>	GUH 160323	<i>Porphyromonas endodontalis</i>	ATCC 35406
<i>Atopobium vaginae</i>	GUH 010535	<i>Porphyromonas gingivalis</i>	ATCC 33277
<i>Bacteroides ureolyticus</i>	GUH 080189	<i>Prevotella bivia</i>	GUH 450429
<i>Bifidobacterium biavatii</i>	GUH 071026	<i>Prevotella disiens</i>	GUH 190184
<i>Bifidobacterium bifidum</i>	GUH 070962	<i>Prevotella intermedia</i>	ATCC 25611
<i>Bifidobacterium breve</i>	GUH 080484	<i>Prevotella melaninogenica</i>	ATCC 25845
<i>Bifidobacterium longum</i>	GUH 180689	<i>Propionibacterium acnes</i> (type I+II)	ATCC 11727/2
<i>Campylobacter gracilis</i>	ATCC 33236	<i>Proteus mirabilis</i>	GUH 07092
<i>Campylobacter rectus</i>	ATCC 33286	<i>Pseudomonas aeruginosa</i>	DSMZ 50071
<i>Campylobacter showae</i>	ATCC 51146	<i>Selenomonas noxia</i>	ATCC 43541
<i>Capnocytophaga gingivalis</i>	ATCC 33612	<i>Staphylococcus anaerobius</i>	DSMZ 20714
<i>Capnocytophaga ochraceae</i>	ATCC 335945	<i>Staphylococcus aureus</i>	ATCC 25923
<i>Capnocytophaga sputigena</i>	ASTCC 33612	<i>Staphylococcus aureus</i> (yellow)	GUH 070921
<i>Corynebacterium nigricans</i>	GUH 450453	<i>Staphylococcus aureus</i> (white)	GUH 070922
<i>Corynebacterium aurimucosum</i>	GUH 071035	<i>Staphylococcus epidermidis</i>	GUH 130381
<i>Dialister</i> sp.	GUH 071045	<i>Staphylococcus haemolyticus</i>	DSMZ 20263
<i>Escherichia coli</i>	GUH 070903	<i>Streptococcus agalactiae</i>	GUH 230282
<i>Eikenella corrodens</i>	ATCC 23834	<i>Streptococcus anginosus</i>	ATCC 33397
<i>Enterococcus faecalis</i>	GUH 170812	<i>Streptococcus constellatus</i>	ATCC 27823
<i>Enterococcus faecalis</i>	ATCC 29212	<i>Streptococcus gordonii</i>	ATCC 10558
<i>Fusobacterium nucleatum nucleatum</i>	ATCC 25586	<i>Streptococcus intermedius</i>	ATCC 27335
<i>Fusobacterium nucleatum polymorphum</i>	ATCC 10953	<i>Streptococcus mitis</i>	ATCC 49456
<i>Fusobacterium nucleatum naviforme</i>	ATCC 49256	<i>Streptococcus oralis</i>	ATCC 35037
<i>Fusobacterium periodonticum</i>	ATCC 33693	<i>Streptococcus pneumoniae</i>	DSMZ 11866
<i>Gardnerella vaginalis</i>	GUH 080585	<i>Streptococcus sanguinis</i>	ATCC 10556
<i>Haemophilus influenzae</i>	ATCC 49247	<i>Streptococcus mutans</i>	ATCC 25175
<i>Helicobacter pylori</i>	ATCC 43504	<i>Tannerella forsythia</i>	ATCC 43037
<i>Lactobacillus acidophilus</i>	ATCC 11975	<i>Treponema denticola</i>	ATCC 35405
<i>Lactobacillus crispatus</i>	GUH 160342	<i>Treponema socranskii</i>	D40DR2
<i>Lactobacillus gasseri</i>	GUH 17085	<i>Varibaculum cambriense</i>	GUH 070917
<i>Lactobacillus iners</i>	GUH 160334	<i>Veillonella parvula</i>	ATCC 10790

ATCC = American Type Culture Collection; D: sample from Forsyth Institute, Boston, MA; GUH = Ghent University Hospital Collection, Ghent, Belgium.

### Microbiological processing

The microbiological samples were frozen at  $-20^{\circ}\text{C}$ . Samples were sent to the oral microbiology laboratory at the University of Bern, Switzerland, and then processed within 3 months. The swabs were transferred to deoxyribonuclease free laboratory tubes (natural flat cap microcentrifuge tubes, Starlab GmbH Ahrensburg, Germany) with 350  $\mu\text{L}$  Tris EDTA buffer (10 mM Tris-HCl, 1.0 mM EDTA and pH 7.6). The swabs were carefully rotated in the buffer solution and then squeezed against the tube walls to recover as much bacterial material in the solution as possible. The swab was then removed. Subsequently, 200  $\mu\text{L}$  of freshly made .5 M NaOH was then added to each vial. The content was then aliquoted in two equal portions in laboratory tubes. Samples were processed as described for the checkerboard DNA-DNA hybridization method described elsewhere [21–24]. The tubes of collected paper points were sonicated for 20 seconds and the paper points were removed. The remaining content was pipetted on to slots and processed as described for the checkerboard DNA-DNA hybridization method [21–24]. Information on the species used in the present study for the checkerboard method is listed (Table 1). Signals were detected by chemiluminescence using the Storm Fluor-Imager (Storm 840, Amersham Bio-

sciences, Piscataway, NJ) with a setup of 200  $\mu\text{m}$  and 600 V. The digitized information was analyzed by a software program (ImageQuant, Amersham Pharmacia, Piscataway, NJ) allowing comparison of the density 19 sample-lanes against the two standard-lanes ( $10^5$  or  $10^6$  cells) and converted to absolute counts by comparisons with these standards. Relative microbial counts were used when different sampling sites (piercing channel, stud, tongue) were compared. The surface area of each of the test studs was defined (circumference  $\times$  length) and used for normalization of microbiological data (cells/ $\text{mm}^2$ ) to compare biofilms on different piercing materials.

### Statistical analysis

Wilcoxon signed-ranks test,  $\chi^2$  test, and Mann-Whitney test were used for statistical analysis of microbiological findings. Adjustment for multiple comparisons was made and a statistically significant difference was defined by  $p < .001$ . For clinical data, standard descriptive statistics were used to summarize the variables studied. Unless otherwise stated, results are expressed as mean  $\pm$  standard deviation (SD). Variations in demographic and clinical data between groups were assessed by  $\chi^2$  and Kruskal-Wallis test. The data analysis was performed with a

**Table 2**  
Subjects' demographic background and clinical data

	Stainless steel (n = 20)	Titanium (n = 20)	Polypropylene (n = 20)	PTFE (n = 20)
Mean age $\pm$ SD	22.7 $\pm$ 3.7	23.4 $\pm$ 4.6	20.9 $\pm$ 7.6	20.8 $\pm$ 6.8
Gender				
Male, n (%)	2 (10)	3 (15)	2 (10)	5 (25)
Female, n (%)	18 (90)	17 (85)	18 (90)	15 (75)
Smokers (life time exposure)				
Nonsmokers, n (%)	6 (30)	8 (40)	6 (30)	5 (25)
Light smokers, n (%)	0 (0)	0 (0)	0 (0)	0 (0)
Moderate/heavy smokers, n (%)	14 (70)	12 (60)	14 (70)	15 (75)
Characteristics of the stud				
Mean time since piercing, months $\pm$ SD	60.1 $\pm$ 50.9	41.5 $\pm$ 32.4*	59.4 $\pm$ 36.3	73.8 $\pm$ 31
Length of the stud <sup>a</sup> , mm $\pm$ SD	16.4 $\pm$ .8	17.7 $\pm$ 1**	16.4 $\pm$ 4.5	17.2 $\pm$ 3.7
Surface area, mm <sup>2</sup> $\pm$ SD	86.5 $\pm$ 12.3	88.7 $\pm$ 5	82.7 $\pm$ 4.1	89.7 $\pm$ 10.3
Clinical data				
Probing depth, mm $\pm$ SD	1.9 $\pm$ .5	1.9 $\pm$ .6	1.9 $\pm$ .6	2 $\pm$ .4
Clinical attachment level, mm $\pm$ SD	1.9 $\pm$ .6	2 $\pm$ .6	2 $\pm$ .5	2.1 $\pm$ .3
Plaque control record, % $\pm$ SD	39.7 $\pm$ 19.7	34 $\pm$ 18.8	31.9 $\pm$ 20.1	32 $\pm$ 17
BOP, % $\pm$ SD	11.8 $\pm$ 17.8	11.4 $\pm$ 15.3	12.7 $\pm$ 14.8	15.3 $\pm$ 14.3

\*  $p = .03$ .

\*\*  $p = .003$ .

<sup>a</sup> Length of the subject's own stud and the test stud.

statistical software package (SPSS 17.5 for MAC computers, SPSS Inc., Chicago, IL).

## Results

### Subject characteristics

Subject characteristics are presented in Table 2. Five subjects were excluded because of antibiotic medication during the study period. None of the subjects presented with clinical evidence of periodontitis or other exclusionary criteria. A total of 80 subjects (68 women, 12 men) completed the study. All participants were Caucasians, aged 16–36 years (mean age  $\pm$  SD = 22.74  $\pm$  4.47), among whom 31.25% had never smoked, 0% were light smokers (1–912 packs lifetime exposure), and 68.75% were moderate to heavy smokers (>912 packs lifetime exposure). The average time since piercing at examination day was 60.05  $\pm$  38.69 months (range: 6 months–14.25 years, median: 60 months).

### Clinical data

Clinical data are shown in Table 2. No subject presented with localized periodontitis as late complication of the tongue piercing. A total of 23 subjects (28.8%) reported 61 lingual recessions (1.91  $\pm$  .96 mm), which might be late complications of the tongue piercings, with 39 gingival recessions located lingually on lower incisors. Four subjects (5%) reported tooth chipping on one tooth each. All of them could exactly report the situation, about how the tooth chipping occurred because of biting on the tongue piercing. One subject had tooth chipping on six teeth, which could not be related to the piercing. No patient had hyperplastic tissues around the piercing. There was no case with swelling or keloid scarring around the piercing.

### Microbiological analysis

Relative microbial counts showed statistically significant differences between the study locations tongue, piercing channel, and piercing stud (Table 3). Most bacterial species were found at

significantly higher proportions ( $p < .001$ ) in samples from the tongue than from the piercing channel (35/80) and the stud (42/80). On the contrary, of the 80 species, 18 were found at significantly higher proportions in samples from the piercing channel than from the tongue. These included *A actinomycetemcomitans* bY4, *Campylobacter gracilis*, *C rectus*, *C showae*, *Capnocytophaga ochraceae*, *Capnocytophaga sputigena*, *Eubacterium saburreum*, *Fusobacterium nucleatum species nucleatum*, *F nucleatum species polymorphum*, *Leptotrichia buccalis*, *P micra*, *Staphylococcus anaerobius*, *S aureus*, *S haemolyticus*, *Streptococcus anginosus*, *Streptococcus intermedius*, *Streptococcus mutans*, and *Treponema denticola* (Table 3). Additionally, statistical analysis identified significantly higher bacterial proportions ( $p < .001$ ) from studs than from tongue samples for six species. These included the following species: *A actinomycetemcomitans* bY4, *Capnocytophaga gingivalis*, *C gracilis*, *C rectus*, *Propionibacterium acnes*, and *S haemolyticus* (Table 3).

Of those 80 species, eight were presented with significantly higher proportions ( $p < .001$ ) from studs than from piercing channels (*Actinomyces naeslundii* type I and II, *A odontolyticus*, *Eikenella corrodens*, *Escherichia coli*, *Proteus mirabilis*, *Selenomonas noxia*, *Streptococcus sanguinis*, and *Veillonella parvula*) (Table 3).

### Analysis of bacteria identified from piercing studs according to the piercing material

There were no statistically significant differences between groups in relation to age, gender, smoking status, or clinical baseline data (PD, clinical attachment level, plaque control record, and BOP). Statistically significant differences between groups in piercing characteristics were identified (Table 2): in the Ti group, the studs were significantly longer ( $p = .003$ ), but the duration with piercing was significantly lower ( $p = .03$ ) than in the other groups. No statistically significant differences were found between groups when comparing microbial counts from the tongue or the channel.

Comparing normalized microbial counts from the studs (cells/mm<sup>2</sup>) regarding the material identified that the total microbial load was significantly higher ( $p < .001$ ) on SS piercings



**Table 3**

Relative microbial counts of samples from tongue, piercing channel, and piercing studs; median (25. percentile; 75. percentile)

Tongue		Piercing channel		Studs	
Bacterial species	Relative microbial counts (%)	Bacterial species	Relative microbial counts (%)	Bacterial species	Relative microbial counts (%)
<i>V parvula</i>	22.36 (16.97; 27.32)	<i>S haemolyticus</i>	14.6 (8.47; 22.03)	<i>V parvula</i>	21.48 (4.9; 43.95)
<i>P melaninogenica</i>	4.45 (2.88; 6.72)	<i>P micra</i>	6.49 (3.63; 9.80)	<i>S haemolyticus</i>	6.59 (.40; 17.26)
<i>A vaginae</i>	3.84 (1.19; 8.51)	<i>C showae</i>	5.20 (2.58; 8.25)	<i>E corrodens</i>	3.11 (1.54; 8.29)
<i>S pneumoniae</i>	3.73 (2.90; 4.52)	<i>V parvula</i>	4.20 (2.44; 10.06)	<i>S pneumoniae</i>	2.31 (0; 4.72)
<i>S haemolyticus</i>	3.73 (2.20; 4.46)	<i>F periodonticum</i>	3.03 (1.93; 5.13)	<i>S oralis</i>	1.59 (0; 2.59)
<i>S oralis</i>	3.09 (2.38; 3.76)	<i>P melaninogenica</i>	2.46 (1.54; 3.69)	<i>C showae</i>	1.23 (0; 2.99)
<i>P aeruginosa</i>	2.90 (2.08; 3.70)	<i>A a (b) Y4</i>	2.37 (1.46; 3.48)	<i>A a (b) Y4</i>	1.18 (0; 4.17)
<i>F periodonticum</i>	2.55 (1.40; 3.27)	<i>S mutans</i>	2.33 (1.74; 2.94)	<i>N mucosa</i>	1.04 (0; 2.83)
<i>Aodontolyticus*</i>	2.35 (1.74; 2.86)	<i>L buccalis</i>	2.28 (1.47; 2.86)	<i>P melaninogenica</i>	1.03 (0; 10.31)
<i>L gasseri</i>	2.11 (.86; 6.52)	<i>F n sp. nucleatum</i>	2.18 (1.98; 2.76)	<i>P micra</i>	.98 (0; 2.91)
<i>C showae</i>	2.05 (.75; 3.71)	<i>S oralis</i>	1.91 (1.43; 2.94)	<i>P mirabilis</i>	.94 (0; 2.29)
<i>S mitis</i>	1.99 (1.53; 2.72)	<i>C rectus</i>	1.82 (.98; 2.48)	<i>C rectus</i>	.93 (.44; 1.65)
<i>L acidophilus</i>	1.90 (1.56; 2.14)	<i>C gingivalis</i>	1.73 (1.20; 2.79)	<i>L gasseri</i>	.84 (0; 2.01)
<i>E corrodens</i>	1.65 (1.15; 3.99)	<i>S intermedius</i>	1.70 (0; 3.35)	<i>Dialister sp.</i>	.83 (.44; 1.52)
<i>N mucosa</i>	1.62 (.66; 7.64)	<i>C gracilis</i>	1.67 (1.22; 2.54)	<i>F periodonticum</i>	.83 (.35; 2.32)
<i>P micra</i>	1.52 (.75; 2.52)	<i>S aureus</i>	1.64 (.96; 2.24)	<i>S noxia</i>	.79 (0; 1.74)
<i>Dialister sp.</i>	1.40 (.44; 2.98)	<i>F n sp. polymorphum</i>	1.55 (.88; 2.56)	<i>F n sp. naviforme</i>	.75 (0; 1.41)
<i>E coli</i>	1.39 (.59; 2.75)	<i>S anginosus</i>	1.43 (1.11; 1.91)	<i>A vaginae</i>	.73 (0; 1.41)
<i>P mirabilis</i>	1.37 (.81; 3.65)	<i>A a (a)29,523</i>	1.39 (1.10; 1.96)	<i>L jensenii</i>	.69 (0; 1.81)
<i>L jensenii</i>	1.28 (.43; 2.12)	<i>S mitis</i>	1.33 (0; 3.39)	<i>E coli</i>	.68 (.29; 1.55)
<i>A a (a)29523</i>	1.24 (1.05; 1.72)	<i>S anaerobius</i>	1.31 (.69; 1.91)	<i>L buccalis</i>	.68 (0; 1.50)
<i>L iners</i>	1.18 (.59; 2.29)	<i>C sputigena</i>	1.20 (.85; 1.50)	<i>S mitis</i>	.66 (0; 1.37)
<i>S noxia</i>	1.10 (.74; 2.49)	<i>T denticola</i>	1.18 (0; 1.91)	<i>F n sp. polymorphum</i>	.64 (0; 1.45)
<i>L vaginalis</i>	1.05 (.69; 2.62)	<i>E saburreum</i>	1.11 (.13; 1.85)	<i>A naeslundii 1 and 2</i>	.64 (0; 1.68)
<i>E saburreum</i>	1.02 (.29; 1.55)	<i>S constellatus</i>	1.09 (.60; 1.62)	<i>C gingivalis</i>	.61 (0; 1.79)
<i>S sanguinis</i>	.95 (.69; 1.22)	<i>F n sp. naviforme</i>	.99 (.54; 1.39)	<i>G vaginalis</i>	.62 (0; 2.72)
<i>S gordonii</i>	.95 (.67; 1.22)	<i>Dialister sp.</i>	.89 (.11; 1.29)	<i>F n sp. nucleatum</i>	.59 (0; 1.41)
<i>S aureus</i>	.94 (.56; 1.13)	<i>A vaginae</i>	.85 (.49; 1.29)	<i>S sanguinis</i>	.58 (0; 1.17)
<i>A naeslundii 1 and 2</i>	.89 (.75; 1.13)	<i>Peptoniphilus sp.</i>	.80 (0; 1.97)	<i>E saburreum</i>	.55 (0; 1.49)
<i>P intermedia</i>	.88 (.55; 1.27)	<i>L acidophilus</i>	.73 (.42; 1.06)	<i>S aureus</i>	.52 (0; .98)
<i>S mutans</i>	.85 (.47; 1.21)	<i>G vaginalis</i>	.72 (0; 1.56)	<i>S epidermis</i>	.49 (0; 1.30)
<i>H pylori</i>	.84 (.23; 2.19)	<i>L jensenii</i>	.72 (0; 1.79)	<i>S anaerobius</i>	.49 (.17; .73)
<i>S anginosus</i>	.84 (.55; 1.11)	<i>H pylori</i>	.71 (.27; 1.02)	<i>S anginosus</i>	.47 (0; .98)
<i>A a (b)Y4</i>	.81 (.62; .99)	<i>A vaginalis</i>	.71 (.48; 1.26)	<i>L vaginalis</i>	.41 (0; .83)
<i>S anaerobius</i>	.78 (.39; 1.01)	<i>S epidermis</i>	.68 (.07; 1.19)	<i>Aodontolyticus</i>	.41 (.05; .63)
<i>S epidermis</i>	.75 (.49; 1.96)	<i>E corrodens</i>	.64 (.44; .97)	<i>T socranscii</i>	.38 (0; .92)
<i>F n sp. naviforme</i>	.73 (.49; .97)	<i>L gasseri</i>	.63 (0; 1.70)	<i>H pylori</i>	.37 (0; .61)
<i>B ureolyticus</i>	.73 (.27; 1.48)	<i>E faecalis</i>	.62 (.31; 1.14)	<i>C sputigena</i>	.36 (.13; .77)
<i>G vaginalis</i>	.67 (.33; 7.49)	<i>B ureolyticus</i>	.60 (.36; .99)	<i>Strept gordonii</i>	.35 (0; 1.81)
<i>S agalactiae</i>	.66 (.35; 1.13)	<i>P intermedia</i>	.58 (.37; 1.26)	<i>L iners</i>	.34 (0; 1.26)
<i>S intermedius</i>	.66 (.51; .88)	<i>T socranscii</i>	.52 (0; 1.05)	<i>L acidophilus</i>	.31 (0; .55)
<i>C nigricans</i>	.65 (.29; 1.80)	<i>P mirabilis</i>	.52 (0; .85)	<i>S agalactiae</i>	.30 (0; 1.33)
<i>F n sp. polym.</i>	.62 (.42; .87)	<i>S aureus (yellow)</i>	.52 (0; .81)	<i>B longum</i>	.29 (0; .52)
<i>L buccalis</i>	.62 (.46; .83)	<i>P anaerobius</i>	.48 (0; 1.69)	<i>P disiens</i>	.27 (0; 1.12)
<i>A israelii</i>	.61 (.48; .78)	<i>S gordonii</i>	.48 (0; .71)	<i>B ureolyticus</i>	.26 (0; .58)
<i>P bivia</i>	.61 (.24; 1.52)	<i>S agalactiae</i>	.47 (0; .72)	<i>A vaginalis</i>	.25 (0; .63)
<i>F n sp. nucleatum</i>	.58 (.44; .81)	<i>P gingivalis</i>	.45 (0; .79)	<i>P anaerobius</i>	.23 (0; .57)
<i>T socranscii</i>	.58 (.35; 1.54)	<i>P disiens</i>	.43 (0; .73)	<i>H influenza</i>	.23 (0; .54)
<i>B longum</i>	.54 (.32; 8.43)	<i>P bivia</i>	.41 (0; .67)	<i>S aureus (white)</i>	.20 (0; .39)
<i>T forsythia</i>	.52 (.39; .67)	<i>S aureus (white)</i>	.37 (0; .73)	<i>E faecalis</i>	.16 (0; .34)
<i>C gingivalis</i>	.49 (.37; .69)	<i>P acnes</i>	.35 (.18; .59)	<i>P aeruginosa</i>	.16 (0; .35)
<i>S constellatus</i>	.49 (.39; .67)	<i>L iners</i>	.34 (0; .67)	<i>C gracilis</i>	0 (0; 2.65)
<i>P gingivalis</i>	.48 (.41; .63)	<i>L vaginalis</i>	.32 (.16; .45)	<i>P intermedia</i>	0 (0; 1.07)
<i>C rectus</i>	.47 (.38; .89)	<i>H influenza</i>	.27 (0; .48)	<i>S intermedius</i>	0 (0; .76)
<i>E faecalis</i>	.46 (.25; 1.69)	<i>L cispatus</i>	.26 (.15; .42)	<i>S mutans</i>	0 (0; .71)
<i>P disiens</i>	.42 (.18; 2.54)	<i>B longum</i>	.24 (.09; .38)	<i>A israelii</i>	0 (0; .67)
<i>P anaerobius</i>	.41 (.28; .68)	<i>C ochracea</i>	.07 (0; .85)	<i>A a (a) 29,523</i>	0 (0; .54)
<i>A parvulum</i>	.378 (.13; 1.29)	<i>N mucosa</i>	0 (0; 7.45)	<i>S constellatus</i>	0 (0; .48)
<i>L cispatus</i>	.37 (.15; .63)	<i>P aeruginosa</i>	0 (0; 4.08)	<i>P bivia</i>	0 (0; .40)
<i>H influenza</i>	.34 (.16; 1.13)	<i>S pneumoniae</i>	0 (0; 1.51)	<i>P gingivalis</i>	0 (0; .39)
<i>A christensenii</i>	.34 (.15; 1.53)	<i>T forsythia</i>	0 (0; 1.47)	<i>A parvulum</i>	0 (0; .33)
<i>A neuii</i>	.33 (.15; 1.76)	<i>S sanguinis</i>	0 (0; 1.23)	<i>P endodontalis</i>	0 (0; .31)
<i>C pseudogenitalium</i>	.33 (.17; 1.32)	<i>E coli</i>	0 (0; .71)	<i>P acnes</i>	0 (0; .27)
<i>C sputigena</i>	.31 (.25; .45)	<i>A christensenii</i>	0 (0; .66)	<i>C nigricans</i>	0 (0; .26)
<i>C gracilis</i>	.30 (.17; .75)	<i>A naeslundii 1 and 2</i>	0 (0; .55)	<i>Peptoniphilus sp.</i>	0 (0; .25)
<i>S aureus (white)</i>	.30 (.19; 1.47)	<i>S noxia</i>	0 (0; .39)	<i>B breve</i>	0 (0; .21)
<i>M curtisii</i>	.29 (.17; .49)	<i>B biavatii</i>	0 (0; .36)	<i>L cispatus</i>	0 (0; .19)
<i>P acnes*</i>	.29 (.24; .47)	<i>V cambriensis</i>	0 (0; .32)	<i>A neuii</i>	0 (0; .17)
<i>A vaginalis</i>	.26 (.14; 1.47)	<i>P endodontalis</i>	0 (0; .22)	<i>C pseudogenitalium</i>	0 (0; .15)

**Table 3**  
Continued

Tongue		Piercing channel		Studs	
Bacterial species	Relative microbial counts (%)	Bacterial species	Relative microbial counts (%)	Bacterial species	Relative microbial counts (%)
<i>B bifidum</i>	.26 (.14; .52)	<i>A israelii</i>	0 (0; .18)	<i>B biavatii</i>	0 (0; .14)
<i>B breve</i>	.24 (.14; .58)	<i>A odontolyticus</i>	0 (0; 0)	<i>M curtisii</i>	0 (0; .13)
<i>S aureus</i> (yellow)	.23 (.09; 1.31)	<i>A neuui</i>	0 (0; 0)	<i>M mulieris</i>	0 (0; .11)
<i>Peptoniphilus</i> sp.	.22 (.16; 1.14)	<i>A parvulum</i>	0 (0; 0)	<i>T denticola</i>	0 (0; .09)
<i>T denticola</i>	.19 (.14; .32)	<i>B bifidum</i>	0 (0; 0)	<i>C ochracea</i>	0 (0; .04)
<i>M mulieris</i>	.17 (.101; .31)	<i>B breve</i>	0 (0; 0)	<i>T forsythia</i>	0 (0; 0)
<i>B biavatii</i>	.17 (.11; .41)	<i>C nigricans</i>	0 (0; 0)	<i>A christensenii</i>	0 (0; 0)
<i>P endodontalis</i>	.16 (.09; .29)	<i>C pseudogenitalium</i>	0 (0; 0)	<i>B bifidum</i>	0 (0; 0)
<i>V cambriensis</i>	.13 (.08; .23)	<i>M curtisii</i>	0 (0; 0)	<i>S aureus</i> (yellow)	0 (0; 0)
<i>C ochracea</i>	0 (0; .79)	<i>M mulieris</i>	0 (0; 0)	<i>V cambriensis</i>	0 (0; 0)

(median:  $.82 \times 10^4$  cells/mm<sup>2</sup>) than on all other studs (Ti:  $.34 \times 10^4$  cells/mm<sup>2</sup>, PP:  $.09 \times 10^4$  cells/mm<sup>2</sup>, PTFE:  $.02 \times 10^4$  cells/mm<sup>2</sup>). Of the 18 species 13 were found at significantly higher levels ( $p < .001$ ) on SS than on all other materials. These included: *Actinomyces neuui*, *A odontolyticus*, *A israelii*, *A naeslundii* type I and II, *Bifidobacterium breve*, *Corynebacterium sputigena*, *Fusobacterium periodonticum*, *Lactobacillus iners*, *L vaginalis*, *Peptostreptococcus anaerobius*, *Prevotella disiens*, *S noxia*, and *V parvula* (Table 4). Additionally, of the 80 species, 67 were found at significantly higher levels ( $p < .001$ ) in samples from SS than from PTFE and PP (Table 4). Twenty-eight of these species were also found at significantly higher levels ( $p < .001$ ) in samples from Ti than from PTFE and PP (Table 4). There were no statistically significant differences between samples from PTFE and PP.

## Discussion

Previous studies [5,7,25,26] have shown that the occurrence of gingival recession is one of the main effects of the use of tongue piercings, whose prevalence can vary from 19.2% to 55%. The present study demonstrated the occurrence of lingual recession in 28.8%, and a prevalence of tooth chipping in 5%. A higher incidence of tooth fractures (19.2%–26.7%) has been reported in previous studies [5–7]. These reported differences can be attributed to the different methodologies used, and possible co-factors like piercing material, barbell stem length, and time of wear [5]. Additionally, we had a very young study population ( $22.74 \pm 4.47$ ), which might be a bias of the study. Given the fact that the popularity of having piercings in oral tissue parts is more common among young subjects, the authors considered that the present study population represented the sub-population at the greatest risk for complications with piercing elements.

Consistent with the age of the study subjects, the periodontal health was excellent and no subject presented with periodontitis. The extent of gingivitis as defined by BOP was low and supported by the fact that dental plaque deposits were also low, suggesting a good oral hygiene level among the study subjects. This might explain the low prevalence of gingival recessions. Additionally, it seems reasonable that the counts of bacteria associated with periodontitis should be low. Nevertheless, the periopathogenic bacteria *A actinomycetemcomitans* (Y4), *F nucleatum* species, *P micra*, and *T denticola* were found at significantly higher proportions ( $p < .001$ ) in samples from the piercing channel than from the tongue. This is in accordance with Ziebolz et al [4], who collected microbiological samples from the surface of 12

tongue piercings and analyzed them for the presence of 11 periodontopathogenic bacteria. Therefore, in subjects with periodontitis, the piercing might provide an additional reservoir for periopathogenic bacteria and should be removed in the course of periodontal treatment.

Recent data suggest that the colonization of bacteria from the back of the tongue seems to change in children and adolescent subjects approaching a microbiota similar to that in adults, and with the presence of bacteria associated with periodontitis [27]. Similar bacteria can be identified from bacterial samples obtained from the back of the tongue and in periodontal pockets (sulci) [28]. Such studies have focused on the 40 bacterial species assessed with the checkerboard hybridization assay [22]. In the present study, the assay was expanded to include other bacteria that are known to be associated with biofilm development on foreign metallic and plastic surfaces.

The dominating bacteria from the tongue samples did not include species associated with periodontal disease. As expected, we found not only *Streptococcus oralis*, *V parvula*, and *Neisseria mucosa*, but also *Streptococcus pneumoniae*, *S haemolyticus*, and *Paeruginosa*. *S pneumoniae* is commonly found in the upper respiratory tract. *S haemolyticus* and *P aeruginosa* could potentially be important pathogens in biofilm development on foreign materials inserted in the oral cavity, such as tongue and lip piercing devices, or dental implants [29,30], and in association with periodontitis [31]. *Staphylococci* sp. have been associated with infections at dental Ti implants [32–34].

Of the 80 species, 67 were found at significantly higher levels ( $p < .001$ ) in samples from SS than from PTFE and PP. These included periodontopathogenic bacteria like *Porphyromonas gingivalis*, *Prevotella intermedia*, *F nucleatum* sp., *C rectus* and *P micra*, as well as bacteria associated with systemic infections (e.g., *S aureus*, *E corrodens*, alpha- and beta-hemolytic *Streptococci*, *Enterococcus faecalis*, *Haemophilus influenzae*, and *P aeruginosa*). Also for local piercing infections, the most commonly found causal agents are *S aureus* and *Pseudomonas* sp. [35].

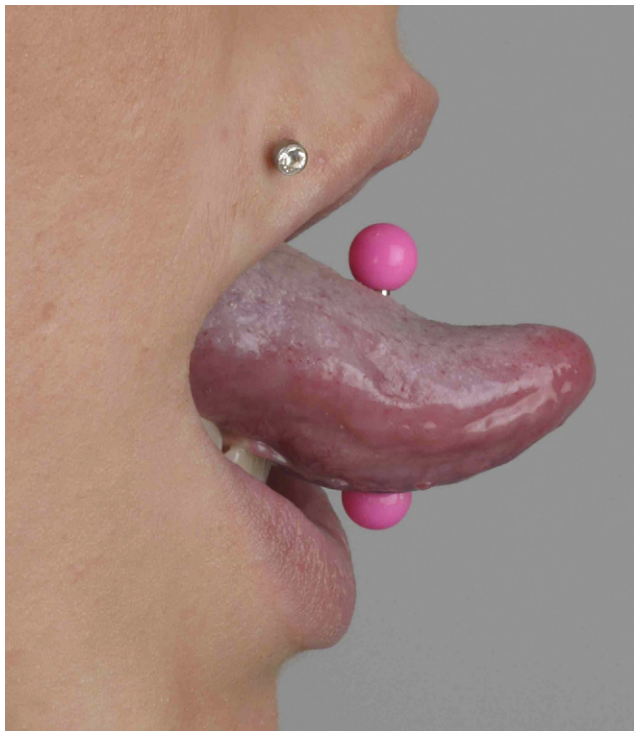
In conclusion, the present study demonstrated low bacterial counts at studs and in piercing channels. Different groups of bacteria are found at higher counts on studs and piercing channels. We confirmed that differences in bacterial colonization patterns occur for different stud materials. Studs made of SS might promote the development of a biofilm, whereas studs made of PTFE and PP may be rather inert to bacterial colonization. The finding of *Staphylococci* on SS and Ti studs may

**Table 4**Distribution of the 67 bacterial species found at significantly higher levels ( $p < .001$ ) in samples from stainless steel than from PTFE and PP; median (25. percentile; 75. percentile)

Bacterial species	Stainless steel counts $\times 10^7$	Titanium counts $\times 10^7$	Polypropylene counts $\times 10^7$	PTFE counts $\times 10^7$
<i>A israelii</i> <sup>ab</sup>	3.98 (2.98; 6.19)	1.23 (0; 2.14)	0 (0; 0)	0 (0; 0)
<i>A naeslundii 1 and 2</i> <sup>ab</sup>	12.56 (7.51; 17.40)	3.41 (1.71; 7.72)	0 (0; 0)	0 (0; 1.19)
<i>A neuii</i> <sup>b</sup>	1.61 (.81; 2.14)	0 (0; .41)	0 (0; .20)	0 (0; 0)
<i>A odontolyticus</i> <sup>ab</sup>	6.55 (5.13; 11.33)	2.21 (1.59; 4.91)	0 (0; .61)	0 (0; 0)
<i>A parvulum</i>	2.92 (1.82; 4.63)	0 (0; 2.14)	0 (0; 0)	0 (0; 0)
<i>A vaginae</i>	6.40 (4.76; 14.78)	4.68 (2.12; 2.14)	0 (0; 2.54)	0 (0; 0)
<i>B biavatii</i>	.99 (.68; 1.69)	.18 (0; 2.14)	0 (0; .54)	0 (0; 0)
<i>B bifidum</i>	1.40 (0; 2.56)	0 (0; 0)	0 (0; 0)	0 (0; 0)
<i>B breve</i> <sup>b</sup>	2.17 (1.12; 2.63)	.51 (0; .79)	0 (0; 0)	0 (0; 0)
<i>B longum</i>	5.47 (1.99; 10.04)	1.23 (.69; 2.95)	0 (0; 1.35)	0 (0; .49)
<i>B ureolyticus</i>	2.24 (1.12; 5.56)	1.22 (.5; 2.14)	.40 (0; .80)	0 (0; .47)
<i>C gingivalis</i> <sup>a</sup>	12.22 (8.92; 22.97)	6.85 (4.32; 8)	0 (0; 0)	0 (0; 2.43)
<i>C nigricans</i>	2.65 (1.37; 4.64)	0 (0; 1.522)	0 (0; 0)	0 (0; 0)
<i>C pseudogenitalium</i>	1.66 (0; 3.66)	0 (0; .93)	0 (0; 0)	0 (0; 0)
<i>C rectus</i>	6.88 (4.07; 15.42)	3.75 (1.967; 7.04)	.58 (0; 2.59)	0 (0; 1.545)
<i>C showae</i>	13.18 (7.23; 28.49)	8.65 (3.06; 18.42)	0 (0; 4.12)	0 (0; 1.00)
<i>C sputigena</i> <sup>b</sup>	13.06 (4.51; 17.66)	3.92 (1.57; 2.14)	0 (0; 2.278)	0 (0; 0)
<i>Dialister sp.</i>	6.30 (3.42; 11.21)	2.77 (1.46; 10.44)	1.44 (0; 2.49)	.67 (0; 1.87)
<i>E coli</i> <sup>a</sup>	10.04 (5.28; 17.02)	3.60 (2.14; 7.34)	0 (0; 1.37)	0 (0; 0)
<i>E corrodens</i>	19.09 (15.44; 40.09)	13.05 (7.62; 2.14)	1.21 (0; 7.62)	0 (0; 0)
<i>E faecalis</i> <sup>a</sup>	2.42 (1.71; 5.69)	.85 (0; 2.24)	0 (0; .64)	0 (0; 0)
<i>E saburreum</i>	5.16 (3.31; 11.49)	1.71 (.95; 5.14)	.39 (0; 1.04)	0 (0; 1.19)
<i>F n sp. naviforme (vincentii)</i>	8.24 (4.29; 17.07)	3.12 (1.76; 2.14)	.29 (0; 1.70)	0 (0; .49)
<i>F nucleatum sp. nucleatum</i>	8.45 (3.47; 12.12)	3.78 (2.27; 2.14)	0 (0; 1.934)	0 (0; 0)
<i>F n sp. polymorphum</i> <sup>a</sup>	10.69 (8.29; 20.12)	4.14 (1.76; 2.14)	0 (0; 0)	0 (0; 0)
<i>F periodonticum</i> <sup>ab</sup>	11.51 (5.98; 19.58)	4.15 (2.36; 2.14)	0 (0; .87)	0 (0; 0)
<i>G vaginalis</i>	5.91 (3.29; 14.63)	2.33 (1.26; 7.58)	.72 (0; 2.42)	0 (0; 2.22)
<i>H influenza</i> <sup>a</sup>	2.09 (.75; 2.89)	1.10 (.46; 2.14)	.16 (0; .46)	0 (0; .25)
<i>H pylori</i>	2.30 (1.550; 5.71)	1.21 (.81; 2.14)	.48 (0; .98)	0 (0; .62)
<i>L acidophilus</i> <sup>a</sup>	9.01 (6.81; 12.74)	4.45 (1.99; 9.156)	0 (0; 0)	0 (0; 0)
<i>L buccalis</i>	6.73 (4.4; 13.24)	2.98 (1.39; 4.74)	1.57 (0; 2.87)	0 (0; 0)
<i>L cispatus</i> <sup>a</sup>	7.34 (5.53; 13.20)	3.77 (1.65; 2.14)	0 (0; 0)	0 (0; .84)
<i>L gasseri</i> <sup>a</sup>	3.79 (2.70; 7.74)	1.39 (1.05; 2.14)	.70 (.46; 1.51)	0 (0; 0)
<i>L iners</i> <sup>ab</sup>	3.73 (1.57; 5.71)	2.24 (.91; 2.14)	0 (0; .88)	.49 (0; .89)
<i>L vaginalis</i> <sup>ab</sup>	1.06 (.75; 2.41)	.67 (0; 2.14)	.49 (.22; .98)	0 (0; 0)
<i>M curtisii</i>	1.25 (.47; 2.11)	0 (0; .81)	0 (0; 0)	0 (0; 0)
<i>N mucosa</i>	37.63 (15.96; 109.01)	16.79 (0; 45.56)	0 (0; 8.589)	0 (0; 0)
<i>P acnes</i>	1.68 (0; 2.26)	.79 (0; 1.756)	0 (0; 0)	0 (0; 0)
<i>P aeruginosa</i>	14.11 (0; 31.92)	5.56 (0; 13.08)	0 (0; 0)	0 (0; 0)
<i>P bivia</i>	3.67 (1.83; 7.73)	0 (0; 1.61)	0 (0; 0)	0 (0; 0)
<i>P disiens</i>	4.06 (1.93; 13.14)	.91 (0; 2.01)	0 (0; .59)	0 (0; .89)
<i>P endodontalis</i>	1.68 (1.9; 3.39)	0 (0; 1.445)	0 (0; 0)	0 (0; 0)
<i>P gingivalis</i>	2.55 (1.46; 5.53)	.61 (0; 2.04)	0 (0; 0)	0 (0; 0)
<i>P intermedia</i> <sup>a</sup>	5.47 (1.40; 10.76)	.994 (0; 4.48)	0 (0; 0)	0 (0; 0)
<i>P melaninogenica</i> <sup>a</sup>	20.21 (9.12; 36.03)	4.53 (3.27; 17.26)	.72 (0; 1.95)	0 (0; .55)
<i>P micra</i>	13.25 (7.06; 28.14)	6.20 (3.66; 14.92)	0 (0; 2.45)	0 (0; 0)
<i>P mirabilis</i> <sup>a</sup>	9.33 (4.19; 17.01)	4.04 (1.59; 7.29)	.789 (0; 1.45)	.53 (0; 1.29)
<i>Peptoniphilus sp.</i>	1.37 (.55; 2.30)	.26 (0; 1.15)	0 (0; 0)	0 (0; 0)
<i>S oralis</i> <sup>a</sup>	2.30 (1.74; 4.24)	1.25 (.93; 2.14)	.86 (0; 2.67)	0 (0; .37)
<i>S agalactiae</i> <sup>a</sup>	3.98 (2.64; 6.51)	2.49 (0; 3.59)	.39 (.06; .87)	0 (0; 1.44)
<i>S anaerobius</i>	3.77 (3.09; 4.51)	2.35 (1.2; 2.14)	0 (0; 1.58)	0 (0; .93)
<i>S anginosus</i>	3.99 (2.49; 6.05)	1.79 (0; 2.14)	0 (0; 1.26)	0 (0; .62)
<i>S aureus</i>	1.48 (1.12; 4.79)	.61 (.52; 2.14)	0 (0; 1.42)	0 (0; .29)
<i>S aureus (white)</i> <sup>a</sup>	0 (0; 1.99)	0 (0; 2.15)	0 (0; .46)	0 (0; 0)
<i>S constellatus</i> <sup>a</sup>	3.87 (3.20; 8.31)	1.40 (1.10; 2.14)	0 (0; 0)	0 (0; .62)
<i>S epidermis</i> <sup>a</sup>	5.28 (4.19; 8.03)	6.17 (2.26; 2.14)	.52 (.345; .81)	0 (0; 0)
<i>S gordonii</i> <sup>a</sup>	12.05 (8.86; 27.66)	9.53 (6.32; 2.14)	0 (0; 0)	1.77 (0; 7.96)
<i>S intermedius</i>	6.57 (4.75; 7.43)	4.52 (0; 8.34)	0 (0; 0)	0 (0; .77)
<i>S mitis</i>	2.74 (1.83; 5.64)	1.58 (0; 3.82)	0 (0; 1.85)	0 (0; 0)
<i>S noxia</i> <sup>ab</sup>	11.50 (7.34; 16.56)	6.61 (3.34; 18.32)	0 (0; 1.26)	0 (0; 1.00)
<i>S pneumoniae</i> <sup>a</sup>	13.30 (11.13; 20.64)	6.88 (4.71; 14.59)	0 (0; 2.89)	0 (0; 0)
<i>S sanguinis</i> <sup>a</sup>	5.1 (3.23; 10.91)	5.41 (2.41; 9.934)	0 (0; 1.25)	0 (0; 0)
<i>T forsythia</i>	0 (0; 4.07)	0 (0; .52)	0 (0; 0)	0 (0; 0)
<i>T socranscii</i>	3.39 (2.42; 5.03)	1.52 (.64; 2.14)	.38 (0; .99)	0 (0; .37)
<i>V parvula</i> <sup>a</sup>	134.47 (51.26; 198.23)	34.44 (19.02; 67.38)	5.19 (.59; 9.70)	0 (0; 3.36)

<sup>a</sup> Bacterial species additionally found at significantly higher levels ( $p < .001$ ) in samples from Ti than from PTFE and PP.<sup>b</sup> Bacterial species additionally found at significantly higher levels ( $p < .001$ ) in samples from SS than from Ti.





**Figure 1.** Piercing of the tongue (and the upper lip). For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.

suggest an elevated risk for complication if the piercing channel is infected. **Figure 1.**

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