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The effect of arginine on oral biofilm communities

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SUMMARY

Alkali production by oral bacteria via the arginine deiminase system (ADS) increases the pH of oral biofilms and reduces the risk for development of carious lesions. This study tested the hypothesis that increased availability of arginine in the oral environment through an exogenous source enhances the ADS activity levels in saliva and dental plaque. Saliva and supra-gingival plaque samples were collected from 19 caries-free (CF) individuals (DMFT = 0) and 19 cariesactive (CA) individuals (DMFT 2) before and after treatment, which comprised the use of a fluoride-free toothpaste containing 1.5% arginine, or a regular fluoride-containing toothpaste twice daily for 4 weeks. ADS activity was measured by quantification of ammonia produced from arginine by oral samples at baseline, after washout period, 4 weeks of treatment, and 2 weeks posttreatment. Higher ADS activity levels were observed in plaque samples from CF compared to those of CA individuals (P = 0.048) at baseline. The use of the arginine toothpaste significantly increased ADS activity in plaque of CA individuals (P = 0.026). The plaque microbial profiles of CA treated with the arginine toothpaste showed a shift in bacterial composition to a healthier community, more similar to that of CF individuals. Thus, an anti-caries effect may be expected from arginine-containing formulations due in large part to the enhancement of ADS activity levels and potential favorable modification to the composition of the oral microbiome.

Keywords

arginine; biofilm; plaque; microflora; caries

SUPPORTING INFORMATION

Additional Supporting Information may be found in the online version of this article:

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INTRODUCTION

Alkali production by oral bacteria influences the ecology of microbial biofilms and the chemical balance between tooth minerals by counteracting the effects of acid produced from sugar metabolism on dental plaque pH (Kleinberg, 1967b; Sissons et al., 1985, 1988; Sissons & Cutress, 1988; Dibdin & Dawes, 1998; Dawes & Dibdin, 2001). Consequently, oral alkali production has the potential to be a major endogenous caries-inhibiting factor (Burne & Marquis, 2000). In fact, considerable knowledge derived from previous clinical and laboratory studies (Stephan, 1944; Kleinberg, 1967a, 1978, 2002; Turtola & Luoma, 1972; Peterson et al., 1985; Dibdin & Shellis, 1988; Margolis et al., 1988; Sissons & Cutress, 1988; Imfeld et al., 1995; Van Wuyckhuyse et al., 1995; Dawes & Dibdin, 2001; Shu et al., 2007; Nascimento et al., 2009, 2013; Morou-Bermudez et al., 2011) supports a significant role of alkali production in oral ecology and inhibition of dental caries. For example, the metabolism of alkali-generating substrates, such as urea and arginine, by oral bacteria elicits a rise in environmental pH (Casiano-Colon & Marquis, 1988; Wijeyeweera & Kleinberg, 1989; Curran et al., 1998). Dental plaque of caries-free individuals has higher pH values and elevated ammonia levels, compared to plaque of caries-active individuals; both at rest and following a carbohydrate challenge (Stephan, 1944; Rosen & Weisenstein, 1965; Turtola & Luoma, 1972; Margolis et al., 1988). Clinical studies performed by our group revealed that increased caries risk is associated with reduced alkali-producing capacity of the microbial populations colonizing the oral cavity of adults (Shu et al., 2007; Nascimento et al., 2009), and children (Nascimento et al., 2013).

The protective principle of oral alkali production is now finding its way to the marketplace as oral care products. For example, toothpaste and mints containing arginine (Acevedo *et al.*, 2005, 2008; Kraivaphan *et al.*, 2013) were demonstrated to be highly effective at inhibiting the initiation and progression of dental caries, so the potential for the cost-effective use of this technology to fight caries appears high. The arginine deiminase system (ADS) is perhaps the single most significant pH homeostasis pathway in oral biofilms that influences susceptibility or resistance to dental caries. If arginolysis is to be targeted as an approach to improve oral health, additional information about the impact of delivery of exogenous arginine from oral health care products on microbial ecology is needed. This clinical study tested the hypothesis that increasing the availability of arginine in the oral environment can increase ADS activity levels in plaque and saliva of individuals with different caries experience and favorably influence the composition of the oral microbiome.

METHODS

Study group

A total of 45 adult subjects (24 females and 21 males; mean age of 26.2 ± 7.8 years) were recruited from dental clinics of the College of Dentistry at the University of Florida (UFCD). The subjects were organized into two groups consisting of 24 caries-free (CF) individuals with no clinical evidence of caries experience [decayed, missing and filled teeth (DMFT = 0)]; and 21 caries-active (CA) individuals with at least two active, cavitated and unrestored carious lesions (DT 2), independently of the number of missing and filled teeth (MFT 0). Of the 45 subjects who were enrolled and participated at the baseline visit, 38 (19 CF and 19

CA) complete all of the study visits. Of the seven subjects who did not complete the study, two moved away from the city, four failed to show up for one of the study visits following the baseline visit, and one was excluded from the study due to self report lack of compliance. Consequently, data was collected from 45 subjects at baseline and from 38 subjects at all other study visits. The selection process excluded subjects younger than 18 years of age and those with fewer than 20 teeth. Other exclusion criteria were systemic diseases; treatment with antibiotics, steroids or any medication known to cause dry mouth in the last 3 months; and the presence of removable or fixed dental appliances. Informed consent was obtained from each volunteer under a protocol approved by the University of Florida Health Science Center Institutional Review Board. Participating subjects completed an enrollment questionnaire describing their demographic characteristics, medical history, oral and dietary habits. Dental plaque and calculus indices were recorded visually as previously described (Greene & Vermillion, 1960). The salivary flow rate, resting pH and buffering capacity were measured during the baseline collection visit using the Saliva-check buffer kit (GC America, IL).

Study design

This single blind, randomized controlled trial had a total duration of 7 weeks and included a washout period of 1 week prior to the treatment phase, a treatment phase of 4 weeks, and a post-treatment phase of 2 weeks. During the washout period, subjects were instructed to brush their teeth twice daily for 1 min using the fluoride-containing toothpaste, *Crest Cavity Protection*®, for 1 week. Subjects were randomly assigned to one of the two treatment groups, which consisted of as follows: (Arg) a fluoride-free toothpaste containing 1.5% arginine, and (F) *Crest Cavity Protection*® (1100 ppm F as NaF). During the treatment phase, subjects were instructed to brush their teeth twice daily for 1 min using their assigned toothpaste for 4 weeks. The recommended amount of toothpaste applied to the toothbrush was shown to subjects as being about 1 g, or 2 cm in length. During the post-treatment phase, subjects discontinued the use of their assigned treatment products and continued to use whatever oral hygiene products and methods they typically used prior to the study.

Sample collection

Subjects were asked to refrain from brushing and flossing their teeth, eating, and drinking anything other than water for 12 h prior to each collection visit. Dental plaque and saliva samples were collected at time zero (baseline), after washout period (washout), after the 4 weeks of treatment (treatment), and 2 weeks after end of treatment (post-treatment). Whole unstimulated saliva was collected by asking the subjects to expectorate into a chilled sterile plastic tube. After saliva collection, supragingival plaque was pooled from all smooth surfaces of the dentition using sterile periodontal curettes. Each portion of plaque scraped from tooth surfaces was immediately transferred to and dispersed in sterile micro-centrifuge tubes containing 10 mM sodium phosphate buffer (pH 7.0). The samples were immediately transported on ice to the laboratory to be analyzed or, if necessary, snap-frozen and stored at -80° C until needed for analysis (Nascimento *et al.*, 2009).

Arginine deiminase system activity levels of oral samples

The biochemical assays for determination of the arginolytic capacity of oral samples were performed following an established protocol (Nascimento *et al.*, 2009). Briefly, ADS activity was measured by quantification of the ammonia produced from the incubation of plaque (25 μ l) and saliva samples (5 μ l) in a mixture containing 50 mM arginine-HCl (Sigma-Aldrich Inc., St. Louis, MO) and 0.5 mM Tris-Maleate buffer (pH 6.0) for 90 min at 37°C. The ammonia produced was detected by the Nessler's Reagent (Sigma-Aldrich Inc.) using ammonium sulfate as the standard. Each sample was assayed in triplicate, and controls for background and interference were always included. ADS activity was normalized to protein content and defined as µmoles of ammonia liberated [minute × (mg of protein)]⁻¹. The detection limit of the assay was 0.01 µmoles of ammonia liberated [minute × (mg of protein)]⁻¹. Protein content was determined as described elsewhere (Nascimento *et al.*, 2009).

Quantitative Real-Time PCR (qRT-PCR)

Real-Time PCR was used to quantify selected acid-and alkali-producing organisms in plaque samples collected at baseline and after 4 weeks of treatment. Species-specific oligonucleotides were used to enumerate Streptococcus mutans (gtfB gene), a well-known cariogenic species and one of the most acidogenic species of the MS group; and two arginolytic species, Streptococcus sanguinis (sagP gene) and Streptococcus gordonii (arcA gene) (Nascimento et al., 2009). The proportions of each species were obtained by normalization to total bacterial counts present in the same plaque sample as determined using universal primers for bacterial 16S rRNA genes (Rupf et al., 1999). These oligonucleotide primers were designed using DNA mfold (http://www.bioinfo.rpi.edu/ applications/mfold/old/) and Beacon Designer 2.0 (Premier Biosoft International, Palo Alto, CA), and have been previously validated (Nascimento et al., 2009). DNA was extracted from plaque using the Ultra Clean Microbial DNA Isolation kit (MO BIO Laboratories Inc., Carlsbad, CA) as recommended by the supplier. A total of 10 ng of genomic DNA from each sample was used in every qRTPCR run. qRTPCRs were carried out and analyzed as described elsewhere (Nascimento et al., 2009). Of note, qRT-PCR does not differentiate viable cells from dead cells.

The Human Oral Microbe Identification Microarray (HOMIM)

The HOMIM analyses were performed at the Human Microbe Identification Microarray (MIM) Core facility at the Forsyth Institute (Cambridge, MA). HOMIM is a 16S rRNAbased microarray that allows for the simultaneous detection of about 300 of the most prevalent oral bacterial species, including many that have not yet been isolated and cultured *in vitro* (Paster *et al.*, 2006; Preza *et al.*, 2008, 2009; Colombo *et al.*, 2009). Briefly, DNA was extracted from plaque samples collected at baseline and after 4 weeks of treatment using the Ultra Clean Microbial DNA Isolation kit (MO BIO Laboratories Inc.). The methods for amplification of 16S rRNA genes, labeling of PCR products and microarray hybridization were performed as detailed elsewhere (Preza *et al.*, 2008; Colombo *et al.*, 2009). The HOMIM microarrays contain 456 unique oligonucleotide probes printed in duplicate on an array that contains a total of 960 printed spots. Image analysis allowed the determination of

the presence or absence of a particular microorganism based on specific criteria set for that individual spot. Data was processed at Forsyth Institute to generate microbial profile maps and cluster analyses. The method can detect the presence of organisms representing about 0.1% of the total sample and the limit of detection is about 10^4 cells.

Data analysis

To determine the sample size, power analyses were performed using proc power-SAS based on the method of ANOVA test. The statistical power calculations were based on the data from a preliminary study. The analyses indicated that 45 subjects were required for addressing the aims of this study with a power of 92% ($\alpha = 0.05$). For descriptive analysis, chi-square test or Fisher's exact test was used to analyze the relationships between categorical variables. In addition, paired T-test was used to evaluate the difference of a continuous variable between baseline and after treatment. As each participant was repeatedly measured at different visits, repeated measures general linear model was used to examine the effects of caries groups and treatment on ADS activity levels and proportions of selected organisms, as determined by qRT-PCR. Akaike's information criterion (AIC) was used to evaluate the model, where smaller AIC values suggest a better model. Contrast statement with the multiple comparisons procedure of the Tukey-Kramer HSD test was used to comparison of ADS and number of bacterial taxa. A *P*-value < 0.05 indicates that the predictor is significantly associated with the average change in the relative proportions of bacterial taxa between visits. All data management and statistical analyses were performed using SAS procedures (SAS 9.1.3; SAS Institute Inc., Cary, NC).

Microbial community profiles were generated from image files of scanned HOMIM arrays using a HOMIM online analysis tool (http://bioinformatics.forsyth.org/homim/). The detection of a particular taxon in a sample was determined by the presence of a fluorescent spot for that unique probe. A mean intensity for each taxon was calculated from the hybridization spots of the same probe and the signals were normalized and calculated as described elsewhere (Colombo et al., 2009). To determine how bacterial community composition varied across samples, total hybridization profiles obtained by HOMIM arrays were compared for each sample using correspondence analysis (CoA) in MeV 4.04 (Saeed et al., 2006). Analysis was done on the absolute intensity HOMIM data (frequency of scores from 0 to 5) and the prevalence of each taxon was computed for each subject and averaged within groups. To identify statistically significant differences between two groups at baseline (CA/CF subjects) and treatment (Arg/F), the Wilcoxon rank-sum test was used with the Benjamini-Hochberg correction and the False Discovery Rate (FDR) to account for multiple testing. To identify statistically significant differences between two groups before and after treatment (baseline/Arg in CF subjects, or baseline/Arg in CA subjects), paired T-test was used with the adjusted Bonferroni analysis and the FDR to account for multiple hypotheses. A *P*-value < 0.05 was considered significant. Sorensen index in estimates was used to compare similarity of the community composition between and within different caries groups. ANOVA analysis with Tukey's test (set to 0.05) on Sorensen indices was used to determine if the community composition was different among groups (JMP 8.0.1).

RESULTS

Baseline ADS activity levels

The ADS activity levels of plaque and saliva at baseline were not significantly correlated to subjects demographic characteristics, medical history, oral and dietary habits (P > 0.05). Plaque and saliva ADS activity at baseline were also not significantly correlated to plaque and calculus index, or to salivary flow rate, resting pH and buffering capacity. While significantly higher ADS activity levels were observed in plaque samples from CF (0.32 \pm 0.28) compared to those of CA subjects (0.27 \pm 0.33; P = 0.048), there were no statistically significant difference in the saliva ADS activity between the caries groups at baseline.

The effect of the arginine toothpaste on ammonia production

A high degree of variability in the rate of ammonia production from arginine was evident among subjects within the caries groups, irrespective of the treatment or time of collection. Table 1 presents the ADS activity levels of plaque and saliva collected from the caries groups at baseline and after the use of toothpaste containing either arginine (Arg) or fluoride (F). The effects of Arg and F on ammonia production via the ADS were dependent on the caries group. Although an increased plaque ADS activity was observed among CF subjects treated with either Arg or F, a statistically significant difference in the levels of ADS activity was observed only among those treated with F (P= 0.042). Among the CA subjects, only those making use of Arg had a statistically significant increase in plaque ADS activity (P= 0.026). General linear regression models with repeated measurements revealed that, under the conditions tested in this study, Arg increased ammonia production in the dental plaque of CA individuals.

There were no statistically significant differences in saliva ADS activity levels among the caries groups after the use of the test toothpastes. Also, there were no statistically significant differences in plaque and saliva ADS activity among the caries groups when the different collection visits (baseline, washout, treatment and post-treatment) were compared.

The effect of the arginine toothpaste on microbial communities

The proportions of two arginolytic species, *S. sanguinis* and *S. gordonii*, and the cariogenic species, *S. mutans*, were determined by qRT-PCR in plaque samples at baseline and after treatment (Supplemental Table S1). Multiple regression analyses revealed no correlation between the frequency of these specific bacterial species and the treatment used and/or the caries group.

Human Oral Microbe Identification Microarray was used to compare the plaque microbiome of the different caries groups before and after the use of Arg or F. Collectively, 263 bacterial species and clusters were detected from the plaque samples. Supplemental Fig. S1 shows HOMIM arrays of plaque samples collected from individuals of the different caries groups and Supplemental Fig. S2 illustrates the microbial profile comparison of the bacterial species and clusters detected.

Table 2 presents the number of bacterial taxa detected from plaque of CA and CF before and after the use of Arg or F. At baseline, a significantly higher average number of bacterial taxa was detected in the plaque of CA (99.7 \pm 8.3) compared to those of CF (81.9 \pm 8.2) subjects (*P* = 0.046). Of particular interest, a greater reduction in the numbers of bacterial species was observed in plaque samples of CA subjects treated with Arg compared to those treated with F.

The microbial profile of CA subjects was distinguished from that of CF subjects at baseline by the frequency of four bacterial species and/or clusters (Table 3). When the Wilcoxon rank-sum test with the Benjamini-Hochberg correction was used to account for multiple testing, no significant difference in bacteria taxa was found (adjusted *P*-value > 0.05). The microbial profiles of the caries groups at baseline are also presented in the Supplemental Fig. S3. The most prevalent species/clusters (>60% of all samples) were *Streptococcus* spp. Clusters III (OT 755/758/767/768/Q65) and II (OT 071/755/758/Q59), Fusobacterium nucleatum ss nucleatum and/or F. nucleatum animalis (OT 420/698/AE01), Haemophilus parainfluenzae (OT 718/W79), Veillonella atypica and/or V. parvula (OT 161/524/Q67), Capnocytophaga granulosa and/or sp clone BB167 (OT 325/326/AA89), Streptococcus oralis and/or sp clones C5MLM037 and EK048 (OT 064/707/F46), Leptotrichia buccalis and/or L. goodfellowii and/or Sneathia sanguinegens (OT 563/837/845/AA45), Streptococcus anginosus and/or S. intermedius (OT 543/644/Q62), Campylobacter gracilis (OT 623/Q04), and Campylobacter concisus and/or C. rectus (OT 575/748/X36). The oral taxon designations (OT) were defined and provided in the Human Oral Microbiome Database (HOMD; www.homd.org).

Table 4 shows the four bacterial species and/or clusters that were detected as being significantly different in the plaque of CA and the one bacterial species and/or clusters that was detected as being significantly different in the plaque of CF before and after treatment with Arg. Supplemental Table S2 shows the list of eighteen bacterial species and/or clusters that were significantly different in abundance in plaque of all subjects, independent of the caries group, before and after treatment with Arg.

Correspondence analysis was also used to analyze the HOMIM data. Figs 1 and 2 show the diversity in the plaque microbial profile of the caries groups at baseline and after treatment with Arg. Overall, the microbiomes of CF subjects were more coherent, presenting a more similar community composition, compared to those of CA subjects. Although the Sorensen index analysis (ANOVA and Tukey's test) showed no statistically significant differences between and within groups, the correspondence analysis graphics showed a trend to a shift in the bacterial composition of plaque from CA treated with Arg to a community more similar to that of CF individuals.

DISCUSSION

Notwithstanding the small sample size and relatively brief treatment period of this study, the results provide evidence that an increased availability of arginine in the oral environment through the use of a toothpaste containing 1.5% arginine can induce dental plaque, but not salivary, bacterial arginine deiminase system (ADS) activity. The effect was especially noted

in caries-active individuals, who present a lower ADS activity at baseline compared to caries-free individuals. Therefore, an anti-caries effect may be expected from toothpaste containing arginine due, in large part, to the ability of arginine to serve as an inducer of the ADS and a substrate for ammonia production by plaque bacteria. In addition, toothpaste containing arginine may also enhance the ability of plaque bacteria to produce ammonia from naturally occurring substrates in saliva. This study supports the hypothesis that protection against caries development may be achieved by increasing the alkali-generating capacity of dental plaque (Liu *et al.*, 2012b; Nascimento *et al.*, 2013).

The reduction in the total number of bacterial species detected among the caries groups after treatment with the fluoride-free toothpaste containing arginine and the fluoride-containing toothpaste, al-thought not statistically significant, is likely due to an overall reduction of plaque formation associated with the introduction of a standardized oral care regimen. As previously observed (Nascimento et al., 2009), a high degree of variability in the rate of ammonia production was evident among subjects within the caries groups, irrespective of the treatment or time of collection. Our qRT-PCR results support the hypothesis that the individual oral alkalinogenic capacity may not be directly associated with a simple change in the proportions of the known ADS-positive bacterial species, S. sanguinis and S. gordonii (Nascimento et al., 2009). Instead, other bacterial species may be responsible for the bulk of oral alkali generation. Also, strain heterogeneity, differences in expression levels of the ADS genes, and/or differential inhibition of enzymatic activity may account for the high degree of variability in ammonia production in health and disease. Recently, our research group demonstrated that the microbial basis for intra-subject variations in oral arginolysis is far more complex than previously appreciated; not only may the arginolytic potential of oral biofilms be associated with the carriage of certain strains of bacteria, but also arginolytic species display a range of ADS activity as a function of environmental factors (Liu et al., 2012a). Together, these findings highlight the need for studies to characterize the bacterial species and/or microbial associations capable of contributing to total arginolysis in the oral cavity.

As determined by HOMIM, caries-active individuals presented a distinct microbial profile compared to caries-free individuals at baseline. The use of the fluoride-free toothpaste containing arginine by caries-active individuals promoted a change in plaque bacterial composition to an apparently healthier community, which is comparable to those of caries-free individuals. Here, it was demonstrated that the HOMIM technique significantly supported our goal of characterizing the microbial profile of individuals with different caries status and oral arginolytic capacities. Most importantly, the HOMIM results suggest that the modulation of the alkali-generating potential of dental plaque, perhaps by using arginine formulations, may foster an ecologically healthy oral environment that could have the potential to suppress the emergence of acid-tolerant, caries-associated pathogenic organisms. Two questions still to be answered are whether the change in the proportions of specific bacterial species and/or clusters is associated solely with the use of the fluoride-free toothpaste containing arginine and whether the identified group of species/clusters is indicative of a favorable and healthy change in the microbiome.

Much remains to be investigated with respect to the effects of novel oral care technologies containing arginine and fluoride (Acevedo *et al.*, 2005, 2008; Kraivaphan *et al.*, 2013) on the composition, gene expression profiles and biochemical activities of the oral microbiota. Nonetheless, based on existing knowledge of the impact of the ADS on pH homeostasis and bacterial bioenergetics, it is expected that the combined mechanisms of action of arginine and fluoride may impact the ecology of oral biofilms in a way that promotes dental health. Future studies will also continue to dissect the underlying molecular basis for the individual and caries-related differences in arginolysis. The combination of these studies will provide information about the various microbial profiles and will determine those that may be considered health- or disease-related, ultimately allowing for the identification of those individuals at greater risk for developing dental caries. Such experiments will expand the knowledge on the diversity of the oral alkali-generating bacteria and their role in oral health and disease.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Correspondence analysis of HOMIM data at baseline and after 4 weeks of treatment with the arginine toothpaste.



Figure 2.

Overlaid correspondence analysis of HOMIM data showing the plaque microbiomes of the caries groups before and after treatment with Arg.

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Table 1

Dental plaque and saliva ADS activity levels of caries-free and caries-active individuals before and after 4 weeks of treatment

	Plaque ADS	± SD	Saliva ADS :	± SD
Caries group	Baseline	After treatment	Baseline	After treatment
CF	0.32 ± 0.28	Arg: 0.41 ± 0.40	0.04 ± 0.04	Arg: 0.03 ± 0.03
		F: 0.58 ± 0.34		F: 0.04 ± 0.03
CA	0.27 ± 0.33	Arg: 0.57 ± 0.51	0.03 ± 0.01	Arg: 0.03 ± 0.02
		F: 0.24 ± 0.08		F: 0.04 ± 0.02

Arginine Deiminase System (ADS) activity levels: µmoles of ammonia liberated min⁻¹ (mg protein); SD, standard deviation; CF, caries-free individuals; CA, caries-active individuals; Arg, fluoride-free toothpaste containing 1.5% arginine; and F, fluoride-containing toothpaste.

Table 2

Number of bacterial taxa detected from plaque before and after 4 weeks of treatment as determined by HOMIM

	Number of	species ± SD
Caries group	Baseline	After treatment
CF	81.9 ± 8.2	Arg: 59.5 ± 6.5
		$F{:}~58.3\pm8.5$
CA	99.7 ± 8.3	Arg: 65.5 ± 6.8
		$F{:}\ 81.6\pm8.5$

SD, standard deviation; CF, caries-free individuals; CA, caries-active individuals; Arg, fluoride-free toothpaste containing 1.5% arginine; F, fluoride-containing toothpaste.

Table 3

Mean frequencies and percentages of bacterial species that were significantly different in plaque of caries-free compared to caries-active subjects at baseline

HOMIM Probe ID	n (%) CF	n (%) CA	P-value*	Adj. <i>P</i> -value
Selenomonas artemidis (OT124/AA49)	19 (79%)	7 (33%)	0.0022	0.31
Veillonella atypica (OT524/W88)	18 (75%)	6 (29%)	0.0026	0.31
Prevotella Cluster IV (OT658/693/714/782/AA44)	11 (46%)	1 (5%)	0.0036	0.31
Streptococcus sp. (OT070/071/N20/Hans H6/7A)	16 (67%)	6 (29%)	0.0071	0.45

CF, caries-free subjects; CA, caries-active subjects; OT, Oral taxon designations for each species are defined and provided in the Human Oral Microbiome Database (http://www.homd.org/);

*Wilcoxon rank-sum test with the Benjamini-Hochberg correction to account for multiple testing.

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Table 4

Mean frequencies of bacterial species that were significantly different in plaque of caries-active and caries-free subjects before and after 4 weeks of treatment with the arginine toothpaste

HOMIM Probe ID	Mean CA baseline ± SD	Mean CA Arg ± SD	n (%) CA baseline	<i>n</i> (%) CA Arg	Absolute <i>t</i> -value	Raw P-value	Adj. <i>P</i> -value
Veillonella parvula (OT161/D96)	1.5 ± 1.2	0	11 (58%)	0	5.5	3.66E-05	0.0095
Fusobacterium nucleatum ss nucleatum and animalis (OT420/698/AD99)	1.4 ± 0.9	0	12 (63%)	0	6.2	9.13E-06	0.0024
Prevotella Cluster IV (OT658/693/714/782/AA44)	1.8 ± 0.6	0.6 ± 0.8	17 (90%)	12 (63%)	6.4	6.41E-06	0.0017
Slackia exigua (OT602/AC97)	1.3 ± 1.2	0	17 (90%)	0	6.4	1.26E-04	00328
HOMIM Probe ID	Mean CF baseline \pm SD	Mean CF Arg ± SD	n (%) CF baseline	n (%) CF Arg	Absolute <i>t</i> -value	Raw <i>P</i> -value	Adj. <i>P</i> -value
<i>Kingella</i> oralis and <i>Neisseria</i> sp. clone BM052 (OT009_706_086)	2.4 ± 1.2	1.2 ± 1.1	17 (90%)	13 (68%)	5.9	1.73E-05	0.0045

Microbiome Database (http://www.homd.org); The analyses used data from the subjects who participated on the baseline and 4 weeks of treatment study visits. Paired *F*test *P* < 0.05. Adjusted Bonferroni CA, caries-active subjects; CF, caries-free subjects; Arg, fluoride-free toothpaste containing 1.5% arginine; OT, Oral taxon designations for each species are defined and provided in the Human Oral correction was applied to account for multiple testing. Degrees of freedom = 17; FDR < 0.01.