



Comparative in vitro investigation of the cariogenic potential of bifidobacteria



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ABSTRACT

Objective: This study aimed to assess the in vitro cariogenic potential of some *Bifidobacterium* species in comparison with caries-associated bacteria.

Design: *Bifidobacterium lactis*, *Bifidobacterium longum*, *Bifidobacterium animalis*, *Bifidobacterium dentium*, *Lactobacillus acidophilus*, *Lactobacillus casei*, *Actinomyces israelii*, *Streptococcus sobrinus* and *Streptococcus mutans* were tested for acidogenicity and aciduricity by measuring the pH of the cultures after growth in glucose and bacterial growth after exposure to acid solutions. Biofilm biomass was determined for each species either alone or associated with *S. mutans* or *S. mutans/S. sobrinus*. Enamel hardness was analyzed before and after 7-days biofilm formation using bacterial combinations.

Results: *B. animalis* and *B. longum* were the most acidogenic and aciduric strains, comparable to caries-associated bacteria, such as *S. mutans* and *L. casei*. All species had a significantly increased biofilm when combined either with *S. mutans* or with *S. mutans/S. sobrinus*. The greatest enamel surface loss was produced when *B. longum* or *B. animalis* were inoculated with *S. mutans*, similar to *L. casei* and *S. sobrinus*. All strains induced similar enamel demineralization when combined with *S. mutans/S. sobrinus*, except by *B. lactis*.

Conclusion: The ability to produce acidic environments and to enhance biofilm formation leading to increased demineralization may mean that *Bifidobacterium* species, especially *B. animalis* and *B. longum*, are potentially cariogenic.

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1. Introduction

Dental caries is a biofilm-associated disease highly associated with frequent intake of dietary sugars. Fermentation of these sugars by biofilm microorganisms leads to acid production, which disrupts biofilm microbial homeostasis and can cause dissolution of tooth minerals (Marsh, 2003). The bacteria considered the most cariogenic are the mutans streptococci, especially *Streptococcus mutans* (van Houte, Sansone, Joshipura, & Kent, 1991; Mattos-

Graner, Correa, Latorre, Peres, & Mayer, 2001). Other acidogenic and aciduric bacterial species, including *Actinomyces* spp. and *Lactobacillus* spp., have been observed in dental caries lesion development (Sansone, van Houte, Joshipura, Kent, & Margolis, 1993; van Houte, Lopman, & Kent, 1996). Additionally, *Lactobacillus* and *Bifidobacterium* species have been identified as part of the dental biofilm on white spot lesions (Van Ruyven, Lingstrom, van Houte, & Kent, 2000).

The *Bifidobacteriaceae* family consists of seven genera: *Bifidobacterium*, *Aeriscardovia*, *Falcivibrio*, *Gardnerella*, *Parascardovia*, *Scardovia* and *Alloscardovia*, which have been isolated from both animals and humans. These bacteria are anaerobic, gram-positive, rod-shaped, and principally colonize the gastrointestinal tract. Species in the genus *Bifidobacterium* have recently received

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significant attention in the consumer food industry due to their beneficial roles in human health. They have been shown to have a role in increasing the adaptive immune response, in treating or preventing respiratory and urogenital infections, and in the prevention of allergies and atopic diseases during childhood (Saxelin, Tynkkynen, Mattila-Sandholm, & de Vos, 2005). Several studies suggest that the consumption of products containing probiotics, such as bifidobacteria, could suppress *S. mutans* levels in saliva (Caglar et al., 2005, 2007, 2008; Cildir et al., 2009). However, recent clinical randomized studies have shown that the administration of probiotics did not affect the *mutans* streptococci levels or the occurrence of dental caries in children (Taipale, Pienihäkkinen, Salminen, Jokela, & Söderling, 2012; Taipale, Pienihäkkinen, Alanen, Jokela, & Söderling, 2013; Nozari, Motamedifar, Seifi, Hatamizargaran, & Ranjbar, 2015). A confounding issue is that levels of caries-related microorganisms are not always related to the severity of dental caries (Mattos-Graner, Zelante, Line, & Mayer, 1998), making the beneficial effects of probiotics on the control of caries difficult to elucidate.

Bifidobacterium species have been isolated from dental plaque, saliva and dentinal caries (Modesto, Biavati, & Mattarelli, 2006; Beighton et al., 2008; Mantzourani et al., 2009). *B. dentium*, *B. longum* and *B. breve* were only detected in the oral cavities of children with occlusal caries and not in those of caries-free children (Mantzourani et al., 2009). Those authors suggested that these bacteria might have a role in the progression of occlusal caries lesions. These findings suggest that this family of bacteria may be related to caries development, possibly due to their acidogenicity and acid-tolerance (van Houte, Lopman, & Kent, 1996; Nakajo, Takahashi, & Beighton, 2010). Tanner et al. (2011) reported a high prevalence of the *Bifidobacterium* spp. in addition to *Streptococcus* in children with advanced dental caries and severe early childhood caries. Thus, this study aimed to determine the cariogenic potential of *Bifidobacterium* species in comparison with currently recognized caries-associated bacteria. The null hypotheses formulated was that the cariogenic potential of *Bifidobacterium* species, represented by acidogenicity, aciduricity, and the ability to form biofilms and to induce enamel demineralization, is not different compared with cariogenic potential of caries-associated bacteria.

2. Materials and methods

2.1. Bacterial strains and growth conditions

Assays were performed using the following *Bifidobacterium* species: *B. animalis* (from ACTIVIA®), *B. longum* (ATCC 15707), *B. lactis* (LMG 18905) and *B. dentium* (ATCC 27678); *Lactobacillus* species: *L. acidophilus* (ATCC 4356), *L. casei* (ATCC 393); *Streptococcus* species: *S. mutans* (ATCC 25175 and 3VF2), *Streptococcus sobrinus* (ATCC 27607); *Actinomyces* species: *A. israelii* (ATCC 12102). All ATCC strains were obtained from Oswaldo Cruz Foundation (FIOCRUZ, Rio de Janeiro, RJ, Brazil) or André Tosello Foundation (Campinas, SP, Brazil). *B. animalis* was isolated from the yogurt ACTIVIA® in Transgalactosylated Oligosaccharides-Propionate agar with supplement lithium-muporicina (50 mg/L) (TOS-MUP agar; Merck Millipore, Darmstadt, Germany). *S. mutans* 3VF2 is highly acidogenic clinical strain kindly provided by Dr. Renata de Oliveira Mattos-Graner (FOP-UNICAMP) (Mattos-Graner, Napimoga, Fukushima, Duncan, & Smith, 2004). Reactivation of strains on selective media was done as follow: *S. mutans* and *S. sobrinus* on Mitis Salivarius agar with bacitracin (0.2 U/mL) (Difco), *L. casei* and *L. acidophilus* on Rogosa agar (Difco); *B. animalis*, *B. longum*, *B. lactis* and *B. dentium* on TOS-MUP agar (Merck) and *A. israelii* on *Actinomyces* Garrod medium (HiMedia Laboratories, Mumbai,

India). Plates were incubated anaerobically with Anaerocult system (Merck).

2.2. Acidogenicity tests

Up to 5 colonies of each species were individually transferred from agar plates to Brain Heart Infusion broth (BHI, Difco) and incubated at 37 °C for 24 h. Bacterial cultures were diluted 1:20 in BHI and the bacterial growth was monitored by turbidimetry. When the optical density reached 0.5 (approximately 10⁸–10⁹ CFU/mL) at 550 nm, tubes were centrifuged and the pellet was resuspended in a solution of 50 mM KCl and 1 mM MgCl₂. The suspension was centrifuged and the pellet was resuspended in 20 mM phosphate buffer and incubated at 37 °C for 60 min for starvation. After centrifugation, the pellet was resuspended in a solution of 50 mM KCl and 1 mM MgCl₂, the pH of the solution was adjusted to 7.0 and glucose was added to a final concentration of 55.5 mM. The pH of the suspension was evaluated immediately and 5, 15, 30, 60, 120, 180 and 240 min after the addition of glucose using a glass electrode (Orion 720 A+; Orion Research Inc.) previously calibrated with pH standards of pH 4.0 and 7.0. The area under the curve (AUC) was calculated of pH drop, considering pH 2.8 as a cut-off point, using UTHSCSA ImageTool software, version 3.0. The acidogenicity was expressed as the AUC (cm²). Each analysis was performed in three distinct experiments (Arthur et al., 2011; Belli & Marquis, 1991).

2.3. Aciduricity tests

Similar to acidogenicity tests, 24 h broth cultures were diluted and grew until optical density at 550 nm of 0.5. 3 mL of each culture were individually distributed into tubes, which were centrifuged and resuspended in 0.1 M glycine buffer with a pH of 7.0, pH 5.0 or 2.8. Immediately after resuspension (Time 0) and after 60 min (Time 60), 100 µL of each sample was serially diluted and plated on BHI agar plates which were incubated for 24 h at 37 °C. Colony Forming Units/mL (CFU/mL) were then counted (Arthur et al., 2011; Duarte et al., 2008).

2.4. In vitro biofilm formation

After growing in BHI containing 1% sucrose and reaching an optical density of 0.5 at 550 nm, aliquots of 200 µL of each strain were individually transferred to the wells of 96-well plates (single-species). Additionally, equal volumes of each bacterial culture were also combined (at the same optical density) with *S. mutans* (3VF2) (dual-species) or *S. mutans* (3VF2)/*S. sobrinus* (multi-species) to complete a total of 200 µL per well. Triplicate of samples were incubated at 37 °C for 48 h. Wells containing only the culture medium were used as negative controls. After incubation, the plates were washed by immersion in distilled water three times to remove the non-adherent cells. After a brief drying, 150 µL of aqueous 1% crystal violet was added to each well, and the plates were incubated at room temperature for 30 min. The crystal violet solution was then removed and the plates washed again 3 times. The plates were inverted on paper towels and remained for 2 h at room temperature to dry. The crystal violet dye that stained the biofilm was then solubilized by incubating with 200 µL of ethanol per well for 30 min. Then, 100 µL of the dye in ethanol was transferred to wells of a new plate, and read at absorbance of 575 nm in a microplate spectrophotometer (BioTek Instruments, USA) to quantify the biomass of the biofilm (Mattos-Graner, Jin et al., 2001).

2.5. In vitro initial caries lesion induction

This part of the study was approved by the Ethics Committee of Araçatuba Dental School—UNESP (protocol 197/2013). Bovine permanent central incisors were maintained one month in 2% formaldehyde solution, pH 7.0, for disinfection. Enamel blocks (4 × 4 mm) were prepared from the most flat portion of the labial surface of the crown using a precision saw (IsoMet 1000, Buehler, Lake Bluff, IL, USA) with two diamond disks (series 15HC 11–4243—Diamond. Buehler) separated by a spacer disk (thickness, 4 mm) under refrigeration with distilled water. Dentin was adjusted to obtain parallel surfaces between enamel and dentin (± 2 mm thick), using sandpaper (CARBIMET, Buehler) under low speed and cooling. Then, the blocks were sequentially polished using 600-, 800- and 1200-grade water-cooled silicon carbide paper disks (Buehler) with a final polish using a felt disk (Buehler Polishing Cloth 40–7618) moistened with a 1- μ m diamond polishing suspension (Extec Corp., Enfield, CT, USA) (Vieira et al., 2005). Blocks with a surface hardness (SH) between 320 and 380 KHN were selected and randomized using Excel program (Microsoft Inc.) into study groups. Twenty enamel blocks were used per group, ten of them or the combination of the bacterial species with *Streptococcus mutans* (3FV2) (duals-species) and another ten specimens in combination with *S. mutans* and *S. sobrinus* (multi-species). Control groups of *S. mutans* and *S. mutans/S. sobrinus* were also analyzed. The modified protocol of a previous study (Lima, Motisuki, Spolidorio, & Santos-Pinto, 2005) was used to induce artificial caries lesions. The bovine enamel blocks were completely isolated with a thin layer of nail varnish, except for the external surface (area = 16 mm²) and individually placed in modified artificial caries solution (BHI supplemented with 1% yeast extract, 0.5% glucose, 1% sucrose and 2% of the bacterial culture – 10⁸ cells/mL) for 7 days at 37 °C, exchanging the culture medium every 48 h. The blocks were measured for microhardness of the enamel surface before (SH1) and after demineralization (SH2) by means of five impressions alongside the initial five impressions, at a distance of 100 μ m using a Shimadzu HMV-2000 microhardness tester (Shimadzu Corp., Kyoto, Japan), with a load of 25 g for 10 s. The percentage of surface hardness loss (%SHL = [(SH2 – SH1)/SH1] × 100) was then calculated.

2.6. Statistical analysis

Means (standard deviations) of area under the curve (AUC) of pH values obtained for bacterial strains during pH drop (acidogenicity) were calculated and the comparison among bacterial species were tested using ANOVA and Tukey tests. For aciduricity assays, CFU/mL counts were transformed to log₁₀ (CFU). The means of counts obtained at pH 7.0 T0 and T60 were calculated and

considered as 100% of cell growth. Counts of viable cells at pH 5.0 and pH 2.8 were also performed and the percentage of viable cells was calculated based on counts at pH 7.0 and data was analyzed using ANOVA and Tukey tests. Absorbance values (550 nm) obtained after 48 h of bacterial biofilm formation were compared among the groups of bacteria and between each species combined (dual or multi-species) or single species according to ANOVA and Tukey tests. Percentage of surface hardness loss (SHL) induced by dual-species and multi-species biofilm exposure after 7 days was analyzed using ANOVA and Tukey tests. Virulence factors comparisons among the groups of bacteria (*Streptococcus*, *Lactobacillus* and *Bifidobacterium*) were performed using ANOVA and Tukey (acidogenicity/aciduricity/single-species biofilm) or unpaired *t*-Student tests (dual/multi-species biofilm). Statistical analysis was performed using the program SPSS version 17.1, considering $p < 0.05$ as significant.

3. Results

3.1. Acidogenicity tests

Fig. 1 shows the results of the AUC for the bacteria tested in this study. Statistically lower values of AUC (high acidogenicity) were found for *S. sobrinus*, *S. mutans* (3FV2 and ATCC), *L. casei*, *B. animalis*, *B. longum* and *A. israelii*, which were not statistically different from each other. *B. lactis*, *B. dentium* and *L. acidophilus* did not differ from each other but they had the highest values of AUC (low acidogenicity).

3.2. Aciduricity tests

Fig. 2 shows means/standard deviations of the percentage of bacterial growth at pH 5.0 and pH 2.8 relative to growth at pH 7.0 (considered 100%). Analyzing each bacterial strain separately (upper case letters), no difference on percentage of bacterial growth at pH 5.0 between T0 and T60 was found for all the tested strains. The percentage of viable cells decreased at pH 2.8 for all bacteria, except for *B. animalis*, *B. dentium*, *L. acidophilus* and *L. casei*, comparing T0 with T60. The percentage of viable cells at pH 2.8 was statistically lower when compared to pH 5.0 at T0 for all tested bacteria, except for *B. longum*, *L. casei* and *S. mutans* ATCC. Additionally, the percentage of viable cells was statistically lower at pH 2.8 after T60 in comparison with pH 5.0 T60 for all strains tested, except for *L. casei*.

Considering each pH and time separately (lower case letters) at pH 5.0 and T0 the percentage of viable cells of *S. mutans* (ATCC) was not different compared with *B. animalis*, but it was statistically lower compared with the other strains. At pH 2.8 and T0 the highest percentage of viable cells was observed for *B. longum*, *L.*

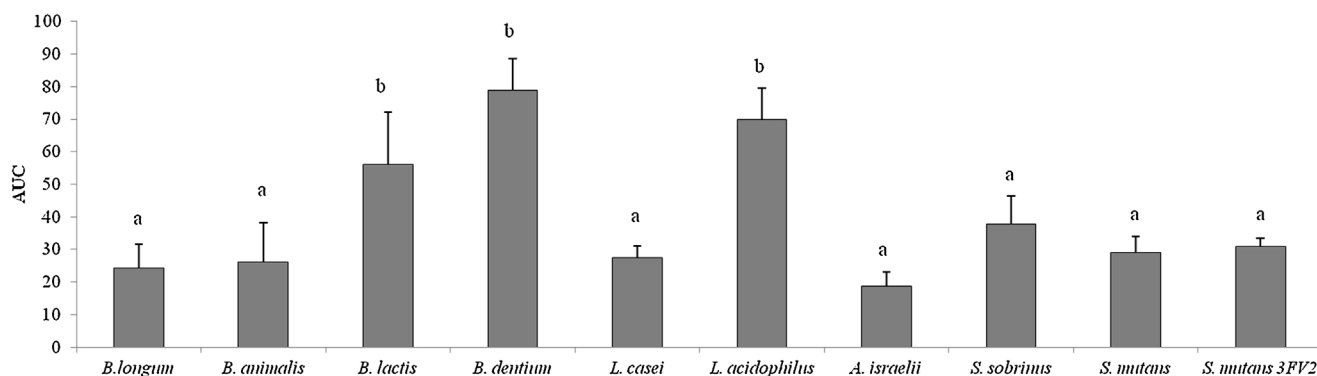


Fig. 1. Acidogenicity (acid production). Means (standard deviations) of area under the curve (AUC) of pH values obtained for bacterial strains during pH drop. ^aDifferent lower case letters show statistical difference among bacterial species, according to ANOVA followed by Tukey tests ($p < 0.05$).

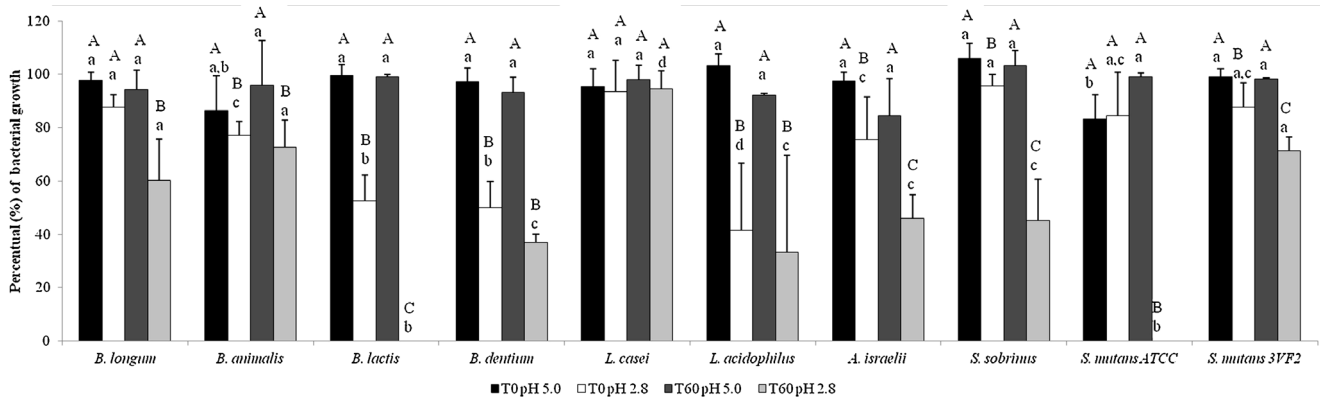


Fig. 2. Aciduricity (acid tolerance). Means (standard deviations) of percentage (%) of bacterial growth obtained after baseline (Time 0) and 60 min (Time 60) of exposition to glycine buffer in pH 5.0 or pH. 2.8 (pH 7.0 was considered as 100%).
^ADifferent upper case letters show statistical difference within each strain tested for both time of exposure (T0 and T60) and pH (5.0 and 2.8), according to ANOVA and Tukey tests ($p < 0.05$).
^aDifferent lower case letters show statistical difference among bacterial strains, considering each pH/time separately (for example *S. mutans* ATCC x *L. casei* in T0 pH5.0), according to ANOVA and Tukey tests ($p < 0.05$).

casei and *S. sobrinus*, but without statistical difference from *S. mutans* (ATCC and 3VF2). At pH 5.0 and T60, no difference was found in the percentage of viable cells among all tested strains. At pH 2.8 and T60, *L. casei* presented the highest aciduricity, followed by *B. longum* and *B. animalis*, followed by *B. dentium*, *L. acidophilus*, *A. israelii*, *S. sobrinus* and *S. mutans* 3VF2, while no growth was observed for *B. lactis* and *S. mutans* (ATCC).

3.3. *In vitro* biofilm formation

Fig. 3 shows means/standard deviations of optical density values (Abs 550 nm) for the bacterial strains evaluated. For single-species biofilms, the ability of *S. sobrinus* to form biofilms was similar to that of *S. mutans* 3VF2, which was higher than biofilms for the other strains. Bifidobacteria and lactobacilli did not form biofilm well. In the dual-species biofilms (associated with *S. mutans* 3VF2), *B. animalis* followed by *L. casei* and *L. acidophilus* had the lowest rates of biofilm formation and *A. israelii* had the highest values, without statistical difference in comparison with *S. sobrinus*

and *B. longum*. In the multi-species biofilm, the association of *S. mutans* 3VF2 + *S. sobrinus* + *B. dentium* presented lower biofilm ability than the combination of *S. mutans* 3VF2 + *S. sobrinus* and *A. israelii* or *L. acidophilus* or *B. longum*, but it was not different compared with the other strains combinations. For all the tested strains, the biofilm formation ability was statistically lower as single-species compared with dual- and multi-species biofilms except for *S. sobrinus* single-species compared with *S. sobrinus* dual-species. Multi-species growth of *B. longum*, *B. animalis*, *L. casei* and *L. acidophilus* presented higher biofilm formation ability compared with their growth as dual-species while no difference was found in biofilm formation ability for *B. lactis*, *B. dentium* and *A. israelii* between dual- or multi-species growth.

3.4. *In vitro* initial caries lesion induction

Fig. 4 shows surface hardness data (%SHL) after exposure of enamel specimens to dual and multi-species biofilms. Under dual-species biofilms, higher demineralization was found in the

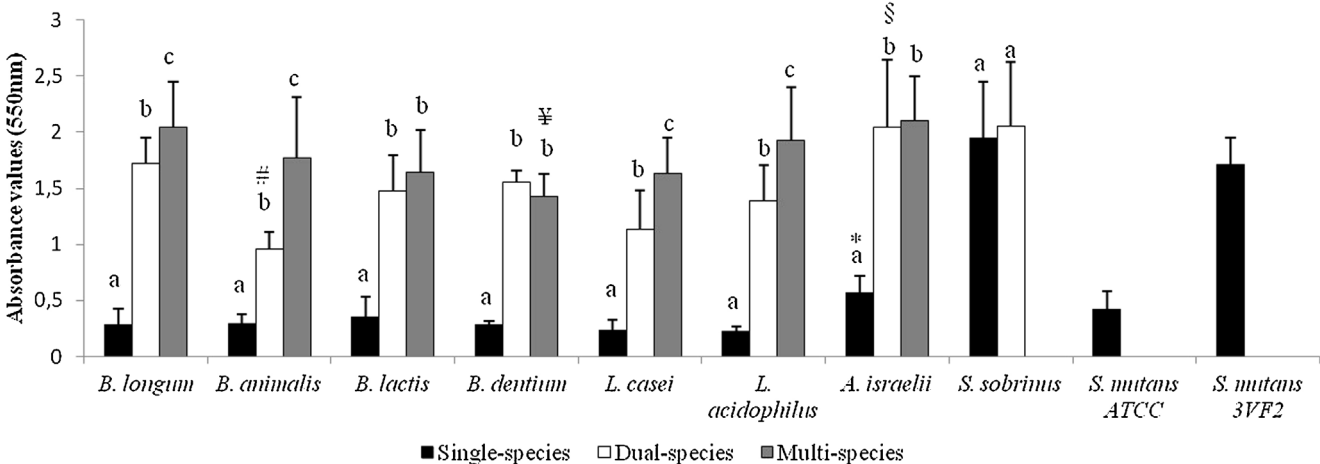


Fig. 3. Biofilm formation. Means (standard deviations) of absorbance values (550 nm) obtained after 48 h of bacterial biofilm formation. *S. sob.*—*S. sobrinus*; 3VF2—*S. mutans* 3VF2.
^aDifferent lower case letters show statistical difference among the biofilm conditions (single, dual or multi-species), considering each species separately, according to ANOVA and Tukey tests.
^{*}Single-species biofilm: Statistical difference between *A. israelii* and other species, according to ANOVA and Tukey tests.
[#]Dual-species biofilm: Statistical difference between *B. animalis* and other species, except by *L. casei* and *L. acidophilus*, according to ANOVA and Tukey tests.
[§]Dual-species biofilm: Statistical difference between *A. israelii* and other species, except by *B. longum*, according to ANOVA and Tukey tests.
[¥]Multi-species biofilm: Statistical difference among *B. dentium* and the following species: *B. longum*, *L. acidophilus* and *A. israelii*, according to ANOVA and Tukey tests.

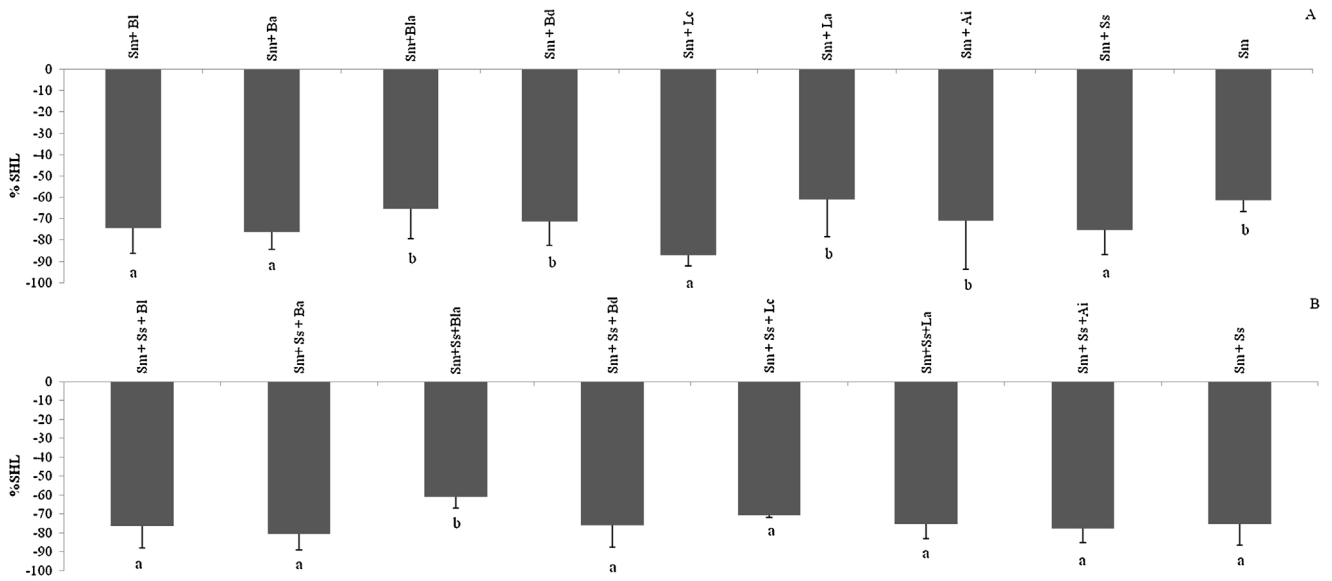


Fig. 4. Enamel demineralization. Percentage of surface hardness loss (%SHL) after 7 days of (A) dual-species and (B) multi-species biofilm exposure. Sm: *S. mutans* 3VF2, Ss: *S. sobrinus*, Bl: *B. longum*, Ba: *B. animalis*, Bla: *B.lactis*, Bd: *B. dentium*, Lc: *L. casei*, La: *L. acidophilus*, Ai: *A. israelii*
^aDifferent letters show statistical difference among the groups, according to ANOVA and Tukey tests.

presence of *B. longum*, *B. animalis*, *L. casei* and *S. sobrinus* while under multi-species biofilms. *B. lactis* associated with *S. mutans* 3VF2 and *S. sobrinus* presented the lowest cariogenic potential compared with the other tested conditions.

3.5. Comparison of the virulence factors among the groups of microorganisms

Table 1 summarizes the comparisons among the groups of bacteria in virulence factors evaluated. *Lactobacillus* and *Streptococcus* species were more aciduric than *Bifidobacterium* species. No difference was found among the groups of microorganisms in acidogenicity. *Lactobacillus* and *Bifidobacterium* species presented similar biofilm formation and induced similar enamel demineralization either in dual-species or in multi-species biofilms.

4. Discussion

Studies have assessed the microbial succession of tooth enamel lesions and noted that various species including *Bifidobacterium* species, in addition to *S. mutans*, may be involved in the development of early caries lesions (Modesto, Biavati, & Mattarelli, 2006; Beighton et al., 2008; Mantzourani et al., 2009; Torlakovic

et al., 2012). The levels of bifidobacteria, streptococci and lactobacilli have been also compared in caries-free and caries-active individuals. Kaur et al. (2013) found that salivary bifidobacterial levels were significantly correlated ($P < 0.001$) with salivary levels of mutans streptococci ($r = 0.732$) and lactobacilli ($r = 0.625$). The frequency of detection of mutans streptococci, lactobacilli and bifidobacteria was significantly higher in caries-active than in the caries-free children, without statistical difference among these bacteria. In adults, salivary levels of bifidobacteria were also significantly related to caries experience and that the salivary levels among mutans streptococci, lactobacilli and bifidobacteria were significantly correlated (Beighton et al., 2010). Specific phenotypic characteristics can influence establishment of species in complex microbial biofilms. The main virulence factors of *S. mutans* that determine their cariogenicity include acidogenicity (production of acids from a variety of fermentable sugars), aciduricity (tolerance to a low pH) and ability to form biofilms on the tooth surface, through adhesion and production of extra and intracellular polysaccharides (Lemos, Abranches, & Burne, 2005).

Data from acidogenicity tests in this current study indicated that, in addition to *S. mutans* (clinical strain), *L. casei*, *B. animalis* and *B. longum* were more acidogenic than the other bacterial

Table 1
 Virulence factors comparisons among the groups of microorganisms.

		<i>Streptococcus</i>	<i>Lactobacillus</i>	<i>Bifidobacterium</i>
Acidogenicity* (AUC)		32.58 (2.13) ^A	52.06(6.74) ^A	46.29 (7.41) ^A
Aciduricity (% bacterial growth)		62.33(7.35) ^A	64.07(5.95) ^A	40.30(5.42) ^B
Biofilm formation (Abs values—550 nm)	Mono-species	1.35(0.03) ^A	0.31(0.03) ^B	0.30(0.02) ^B
	Dual-species	–	1.22(0.05) ^A	1.41(0.06) ^A
	Multi-species	–	1.78(0.07) ^A	1.70(0.06) ^A
Enamel demineralization (%SHL)	Dual-species	–	–69.91(4.44) ^A	–71.75(2.35) ^A
	Multi-species	–	–72.47(1.77) ^A	–73.61(2.40) ^A

^ADifferent upper case letters show statistical difference among the groups. According to ANOVA and Tukey (acidogenicity, aciduricity/mono-species biofilm) or unpaired *t*-Student tests (dual/multi-species biofilm).

* Means (Standard Error of Means).

strains analyzed. *S. sobrinus* and *A. israelii* were among the most acidogenic strains, but had intermediate acid tolerance. The pH drops caused by probiotics, including *B. longum*, *B. lactis*, *L. casei* and *L. acidophilus*, from the degradation of sucrose and other sugars were evaluated by [Haukioja, Loimaranta, & Tenovu \(2008\)](#). The authors found that all of the strains produced acids from glucose with a reduction of pH comparable to that of *S. mutans*, similar to the present results. Those authors also found that all lactobacilli and bifidobacteria tested, except for *L. rhamnosus* GG and *B. lactis* Bb12, caused a significant decrease in pH in the presence of lactose.

The current study showed that the growth of all strains, including species of bifidobacteria, was not affected by pH 5.0 after 60 min of acid exposure. The same was not observed at pH 2.8. Among bifidobacteria, *B. longum* and *B. animalis* were the most acid-resistant strains and *B. dentium* was the least acid-resistant strain, at pH 2.8. These results are in agreement with those obtained by [Nakajo, Takahashi, & Beighton \(2010\)](#) who evaluated the effects of acidification on the survival of *B. dentium* and *B. longum* in comparison with *S. mutans*, *S. sanguinis* and *L. paracasei*. High levels of survival in acidic environments were observed for bifidobacteria, which were comparable to *S. mutans*, demonstrating how these species could co-exist with *Streptococcus mutans* in acidic carious lesions. *B. longum* was the most resistant bacteria at pH 5.0 when compared to the others strains, while *L. paracasei* was the most resistant at pH 4.0. In contrast, *B. dentium* was less resistant to acids among all tested bacteria.

Bacterial aciduricity is maintained by a mechanism in the cell membrane called proton-translocating ATPase (F1F0-ATPase). This mechanism controls the entry of H⁺ ions and maintaining a more basic cytoplasmic pH when compared to the extracellular medium. Furthermore, the F-ATPase system has dual role in the acid tolerance of cells and eliminating protons in certain circumstances it generates ATP (energy) for the growth and persistence of bacterial species ([Lemos & Burne, 2008](#)). Studies have reported that the F-ATPase activity in *Bifidobacterium animalis* and *Bifidobacterium animalis* subsp *lactis* increases with environmental acidification ([Gibson & Wang, 1994](#); [Sánchez, de los Reyes-Gavilán, & Margolles, 2006](#)). [Matsumoto, Ohishi, & Benno \(2004\)](#) evaluated the aciduricity and F-ATPase activity of *B. lactis* (2 strains), *B. animalis* (4 strains), *B. bifidum* (4 strains), *B. breve*, *B. infantis*, *B. catenulatum*, *B. longum*, *B. pseudocatenulatum* and *B. adolescentis* and demonstrated that *B. lactis* and *B. animalis* were able to survive in an acidic environment (they were stable at pH 3–5 for 3 h). They reported that this ability was related to the species and not the strain and was associated with increased F-ATPase activity. In contrast with the current study findings, [Matsumoto et al. \(2004\)](#) reported that *B. longum* was not acid tolerant, with reduced growth by about 3-fold at log₁₀ after exposure to pH 5 for 3 h. Other ATP-independent mechanisms have been studied in *S. mutans*, *S. gordonii*, *S. salivarius* and *L. casei* ([Fozo & Quivey, 2004](#); [Fozo, Kajfasz, & Quivey, 2004](#)). The membrane composition may be altered in the presence of acids, increasing the levels of long-chain monounsaturated fatty acids, allowing the species to become more resistant to acids. The cell membrane of *B. animalis* contains large amounts of these fatty acids ([Ruiz, Sánchez, Ruas-Madiedo, de los Reyes-Gavilán, & Margolles, 2007](#)), suggesting that these bifidobacteria may also display this type of acid tolerance.

In this present study, bifidobacteria and lactobacilli had lower ability to form biofilm as single-species compared with dual- and multi-species biofilms. Multi-species growth of *B. longum*, *B. animalis*, *L. casei* and *L. acidophilus* presented higher biofilm formation ability compared with their growth as dual-species. Previous studies had demonstrated that some bifidobacteria (*B. breve*, *B. longum*, *B. lactis*, *B. adolescentis*, *B. infantis*) exhibited low adhesion to hydroxyapatite discs coated or not coated with saliva

([Haukioja et al., 2006](#); [Nagaoka et al., 2008](#)). This ability was improved when the species had co-adhered with primary colonizers, such as *A. naeslundii*, *V. parvula*, and *F. nucleatum*.

Due their lower capacity to adhere to tooth and to form biofilm by themselves, enamel demineralization was induced only by lactobacilli and bifidobacteria combined with *S. mutans* or with *S. mutans* and *S. sobrinus*. The dual-species biofilms of *S. mutans* with *B. longum*, *L. casei*, *B. animalis* or *S. sobrinus* and the multi-species biofilms of *B. animalis* and *A. israelii* with *S. mutans* and *S. sobrinus* caused the highest loss of surface hardness. Different combinations of *S. mutans* with *L. casei* and *L. acidophilus* were recently tested to observe the induction of non-cavitated enamel lesions in human teeth ([De Campos et al., 2015](#)). Their results showed that dual combination of *S. mutans* and *L. casei* or triple combination of *S. mutans*, *L. casei* and *L. acidophilus* promoted the highest loss in enamel surface hardness. The depth of lesions was analyzed using polarized light microscopy and erosive lesions developed in enamel after 20 days of cariogenic challenge. The results of the present study are consistent with those results, showing that the combination of *L. casei* and *S. mutans* caused a greater enamel demineralization compared to other species tested. Studies evaluating the impact of the biofilm formed by *Bifidobacterium* species and *S. mutans* on demineralization of dental enamel have not yet been found, making comparisons with the present results difficult. It could be concluded that the ability to produce acidic environments and to enhance biofilm formation leading to increased enamel demineralization may mean that *Bifidobacterium* species, especially *B. animalis* and *B. longum*, are potentially cariogenic.

Conflict of interests

The authors declare that there are no conflicts of interest.

Ethical approval

The experimental procedures described here were approved by the Ethics Committee of Araçatuba Dental School – UNESP (protocol 197/2013).

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