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1	Bruno Mello de Matos, Fernanda Lourenção Brighenti, Thuy Do, David Beighton,
2	Cristiane Yumi Koga-Ito
3	
4	Acidogenicity of dual species biofilms of bifidobacteria and Streptococcus mutans
5	
6	Bruno Mello de Matos - UNESP, Univ. Estadual Paulista, São José dos Campos-SP;
7	UNIVASF, Univ. Federal do Vale do São Francisco, Paulo Afonso-BA, Brazil
8	Rua da Alvorada, General Dutra, Paulo Afonso, BA, 48607-190, Brasil
9	Fernanda Lourenção Brighenti - UNESP, Univ. Estadual Paulista, Araraquara-SP,
10	Brazil
11	Rua Humaitá, 1680, Centro, Araraquara, SP, 14801-385, Brasil
12	Thuy Do – School of Dentistry, Faculty of Medicine and Health, University of Leeds,
13	Leeds, UK
14	Division of Oral Biology, Wellcome Trust Brenner Building, Level 7, St James'
15	University Hospital campus, LS9 7TF, Leeds, UK.
16	David Beighton – Dental Institute, King's College London, London, UK
17	Bessemer Rd., Denmark Hill, London SE5 9RW, UK.
18	Cristiane Yumi Koga-Ito – UNESP, Univ. Estadual Paulista, São José dos Campos-SP,
19	Brasil
20	Institute of Science and Technology, UNESP - Avenida Engenheiro Francisco José
21	Longo 777, São Dimas, São José dos Campos, SP, 12245-000, Brazil
22	
23	Corresponding author: Cristiane Yumi Koga Ito, cristiane@fosjc.unesp.br, Telephone:
24	+5512 981253017, Fax: +5512 39479000
25	
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- 34
- 35 Abstract
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Objective: the aim of this study was to evaluate the acidogenicity of dual species 37 38 biofilms of bifidobacteria and Streptococcus mutans. Materials and Methods: the 39 following strains were tested: Bifidobacterium dentium DSM20436, Parascardovia 40 denticolens DSM10105 and Scardovia inopinata DSM10107. Streptococcus mutans UA159 and Lactobacillus acidophilus ATCC4356 were used as control. Bifidobacteria 41 42 were studied planktonically as they were not able to form monospecies biofilm, they were grown in biofilms associated with S. mutans. Endogenous polysaccharide reserves 43 of cultures at log phase were depleted. Standardized suspensions of the microorganisms 44 45 were incubated in growth media supplemented with 10 mM glucose, lactose, raffinose, glucose or xylitol. S. mutans biofilms were grown on glass cover slips for 24 h to which 46 bifidobacteria were added. After 24 h, the dual species biofilms were exposed to the 47 same carbon sources and after 3 h the pH of spent culture media and concentrations of 48 organic acids were measured. Statistical analyses were carried out using ANOVA and 49 Tukey's Test (α =0.05). Results: a higher pH drop was observed when S. mutans was 50 51 associated with P. denticolens or S. inopinata, in either planktonic and biofilm cultures than from S. mutans alone. Bifidobacteria showed a higher pH drop in the presence of 52 53 raffinose than S. mutans or L. acidophilus. Conclusions: Dual species biofilms of bifidobacteria and S. mutans produced more acid and a greater pH drops than biofilms 54 55 of S. mutans alone. Clinical relevance: New insights on the complex process of caries pathogenicity contribute to the establishment of preventive and therapeutic measures, in 56 57 particular in specific cases, such as in early childhood caries.

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59 Keywords: acidogenicity, bifidobacteria, biofilms, dental caries

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64 Introduction

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The etiology of caries is undoubtedly complex. It is generally recognized that 66 microbial, environmental and host factors interact to contribute to dental caries 67 68 development [1]. Although dental caries is a biofilm-mediated disease, it is unlikely that all members of the oral biofilm participate equally in the caries process. The ecological 69 70 plaque hypothesis suggests that the cariogenic oral environment will select increased 71 proportions of acidogenic and aciduric microbiota [2]. These microorganisms include 72 lactobacilli, streptococci, Actinomyces spp., yeasts and bifidobacteria [3]. The resultant 73 pH drop may induce dental enamel demineralization under the critical pH of 5.5 [4].

Aas et al. [5] using molecular techniques demonstrated that 10% of subjects with rampant caries in secondary dentition did not have detectable oral levels of S. mutans in intact enamel and white-spot lesions. The authors suggested that at least half of the bacteria associated with dental caries have not yet been cultivated. Thus, there is a considerable body of evidence for the emergence of other taxa, in addition to S. mutans in a cariogenic oral environment or within carious lesions [3].

Many studies have reported the presence of bifidobacteria in the oral cavity of healthy and diseased children and adults. These bacteria were found in saliva, plaque and dental caries [3, 6-7]. Beighton et al. [8] demonstrated that the bifidobacteria levels in adults' saliva were not significantly different from the levels of mutans streptococci. Similar observations were described in caries active children [9].

Bifidobacterium dentium is the most prevalent bifidobacterial species in the oral cavity [3] with Parascardovia denticolens and Scardovia inopinata also frequently isolated [6, 10].

Bifidobacteriaceae consists of seven genera (Aeriscardovia, Alloscardovia, Bifidobacterium, Falcivibrio, Gardnerella, Parascardovia, Scardovia) and about 36 species, the majority of which have been described and isolated from the intestinal and caecal microbiota. The range of taxa reported to be oral commensal seems primarily restricted to Bifidobacterium dentium, Parascardovia denticolens and Scardovia inopinata [3].

94 Considering that little is known about the influence of environmental factors 95 dietary components in particular - on acid production by oral bifidobacteria, and that

acid production from carbon sources is an important cariogenic feature, the aim of this
study was to evaluate the acidogenicity of bifidobacteria after exposure to different
carbon sources, and determine if bifidobacteria are able to increase the acidogenicity in
single and dual species planktonic cultures or biofilms in association with Streptococcus
mutans.

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102 Materials and Methods

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104 Strains and incubation conditions

105 The following type strains of bifidobacteria were used: Bifidobacterium dentium 106 DSM 20436, Parascardovia denticolens DSM 10105 and Scardovia inopinata DSM 107 10107. Also, Streptococcus mutans UA 159 and Lactobacillus acidophilus ATCC 4356 108 were included. S. mutans is considered an important species related to dental caries 109 initiation whilst lactobacilli are related to dental caries progression.

110 Cultures were obtained for each species from two independent frozen stocks. S. 111 mutans and B. dentium were grown in semi-defined medium broth supplemented with 112 yeast extract (SDMY) and 0.2% sucrose [11]. L. acidophilus, P. denticolens and S. 113 inopinata were grown in Lactobacilli MRS Broth (Difco, USA) developed by De Man, 114 Rogosa and Sharpe [12]. All strains were grown to log phase at 37 °C in anaerobic jars 115 (10% CO₂, 10% H₂, 80% N₂).

Strains were studied planktonically and in biofilms and all experiments wereperformed in duplicate on two different occasions.

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119 Preparation of planktonic cultures

Bacterial cultures were washed twice in cysteine peptone water (CPW) [5 g/l yeast extract, 1 g/l peptone, 8.5 g/l NaCl, 0.5 g/l L-cysteine-HCl]. Depletion of endogenous carbohydrate reserves in stationary phase cultures was performed by incubating the washed cells for 30 min in water bath at 37 °C.

124 Standardized inocula of the microorganisms at $OD_{620}=0.7$ were prepared in an 125 artificial saliva medium modified by McBain et al., 2005 [13], pH 7.0, for pH drop 126 evaluation. 127 Pilot tests showed that, even though McBain medium is an artificial saliva medium, it causes undesirable interferences in organic acids analyses by capillary 128 129 electrophoresis. During those tests, it was verified that the chemical composition of 130 McBain medium produced peaks (observed on Millenium Chromatography Manager 131 Software) that overlap to the acid peaks making them difficult to identify. For this 132 reason, SDM broth (modified version of SDMY broth prepared without yeast extract), 133 supplemented with aqueous solutions of glucose, lactose, raffinose, sucrose or xylitol in final concentrations of 10mM was used to organic acids analyses. Sterile distilled water 134 135 was used as negative control.

136 Dual species suspensions were prepared using S. mutans inocula plus137 bifidobacteria at a 1:1 ratio.

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139 Preparation of biofilms

Pilot studies showed that none of the bifidobacteria species and L. acidophilus was able to form single species biofilms in the model used in the present study. It was therefore not possible to assess the production of acid or utilization of carbohydrates by single species biofilms composed of bifidobacteria or lactobacilli alone. Instead, we assessed the acidogenicity of dual species biofilms formed inoculating either bifidobacteria or lactobacilli onto pre-formed S. mutans biofilms.

To produce the pre-formed S. mutans biofilms, S. mutans cultures were standardized in SDMY ($OD_{620}=0.7$) and diluted 1:50 in SDMY plus 0.2% sucrose. S. mutans biofilms were grown on glass coverslips (\emptyset 12 mm) using an active attachment model [14]. 24-wells plates were filled with 1.5 mL of the diluted inocula per well, and incubated for 24 h. The glass coverslips containing S. mutans biofilms were placed in 24-well plates that were filled with 1.5 mL of either bifidobacteria or L. acidophilus diluted inocula.

The bifidobacteria (B. dentium, P. denticolens or S. inopinata) or L. acidophilus cultures to be added to the S. mutans biofilms were prepared ($OD_{620}=0.7$) and diluted 1:50. Inocula of B. dentium were prepared in SDMY plus 0.2% sucrose and P. denticolens, S. inopinata and L. acidophilus were prepared in Lactobacilli MRS Broth (Difco, USA).

160

After 24 h of incubation, dual species biofilms were washed twice in CPW. Depletion of endogenous polysaccharide reserves was performed by incubating the biofilms in CPW for 30 minutes in water bath at 37 °C.

161 The depleted biofilms were placed in 1.5 ml McBain medium or SDMY broth 162 supplemented with 10 mM glucose, lactose, raffinose, sucrose or xylitol and incubated 163 at 37 °C for 3 h.

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165 Assessment of biofilm viability

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The presence of S. mutans, lactobacilli and bifidobacterial species in mixed 167 168 biofilms was evaluated by culture method. Glass coverslips with biofilms were carefully 169 detached from the clamps and placed in 2 mL CPW. Biofilms were dispersed by 170 sonication on ice for 120 times one second at amplitude of 40 W (Vibra CellTM, Sonics and Materials Inc., Newtown, USA) [15]. Serially diluted samples were plated onto 171 172 SB20 (sucrose bacitracin), Rogosa (Difco, USA) and MMTPY agar plates (modified version of mupirocin trypticase peptone yeast extract) for isolation of S. mutans, 173 lactobacilli and oral bifidobacteria, respectively [8,16]. The plates were incubated 174 175 anaerobically (as previously described) at 37 °C for 48 h. Colonies were counted and 176 expressed as colony forming units (CFU). Experiments were performed in sextuplicate on two different occasions. 177

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180 Assessment of suspensions and biofilms acidogenicity

The pH of the culture medium was measured to estimate biofilm acidogenicity at
0 and 3 h and pH variations calculated. The measurements were performed with the aid
of an electrode with a micro-bulb (Hanna, Woonsocket, Rhode Island, USA).

The amount of organic acids was analyzed by capillary electrophoresis (Waters Capillary Ion Analyzer; Milford, MA, USA) in plates with SDM broth. Samples were run in duplicate, and Millenium Chromatography Manager Software, version 3.05 was used for data analysis. Peak identification and peak area integration were manually corrected if necessary. Sodium salts of formic, acetic, propionic, butyric, succinic, and lactic acid were used to prepare single and mixed standard solutions in deionized water, ranging from 0.05 to 2 mM. Calibration curves were made for each acid separately. As
an internal standard, 0.1 mM oxalic acid was included in all samples. Lactic, propionic,
acetic, formic, butyric, and succinic acid concentrations were determined [17].
Experiments were performed in quadruplicate on two different occasions.

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195 Data analysis

Initially all data were compared to the appropriate water, no added carbohydrate control. Then for those cultures in which significant changes to pH or to acid levels occurred the results obtained for bifidobacteria species were compared to control microorganisms (S. mutans or L. acidophilus). Statistical analysis were carried out using Graphpad Prism 3 (ANOVA and Tukey's Test, α =0.05).

- 201
- 202 Results
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None of the bacteria in the planktonic phase or in biofilms produced significant changes to the pH of the media or to the concentrations of lactic or acetic acids when incubated with xylitol (data not shown). Table 1 shows the pH drop (Δ pH) after the addition of different carbon sources to single species suspensions. For L. acidophilus, the pH drop was higher than S. mutans when glucose was used. Statistically significant Δ pH were observed, which indicate that the presence of raffinose seems to be better metabolized by the three bifidobacterial species than by S. mutans or L. acidophilus.

Higher pH drop was observed when S. mutans was associated with P. denticolens or S. inopinata, in either planktonic or biofilms cultures (Tables 1 and 2). The association between S. mutans and B. dentium in suspension promoted higher pH drop when lactose, raffinose or sucrose was used in comparison to S. mutans or S. mutans and L. acidophilus (Table 1). However, the co-culture of S. mutans and B. dentium in biofilms promoted lower pH drop than S. mutans single species biofilms or S. mutans and L. acidophilus biofilms.

Final pH for enamel demineralisation was below critical (5.5) for all microorganisms and associations when culture media was supplemented with glucose or sucrose. Co-culture of S. mutans/P. denticolens and S. mutans/S. inopinata led to the highest pH drop in presence of glucose (minimum final pH 4.3), sucrose (minimum final pH 4.2), and raffinose (minimum final pH 4.6). The pH (5.5) was not reached in
the presence of lactose, xylitol or control. Raffinose promoted a pH drop below critical
pH in either single species suspensions or associated to other species, both in planktonic
and biofilms. For this carbohydrate, final pH from S. mutans or S. mutans and L.
acidophilus cultures remained above critical levels (pH 6.2 and 6.3, respectively)
(supplementary material).

Table 3 displays the organic acids production of single species and dual species suspensions. Butyric, formic, propionic and succinic acids were below detection limit (0.01 mM, according to Kara et al., 2006 [17]). For B. dentium in single species cultures, more lactate is produced in the presence of raffinose or sucrose. When associated to S. mutans, lactate production was higher in the presence of raffinose. P. denticolens produces more acetate for all carbohydrates.

The same pattern of lactate production is observed for either single species or dual species suspensions of S. inopinata. On the other hand, while acetate production in S. inopinata single species is higher than S. mutans for glucose, raffinose and sucrose, in dual species cultures, significantly higher concentrations of acetate was found for all carbon sources.

239 In dual-species biofilms, the combination of S. mutans and B. dentium did not produce more acid than S. mutans or S. mutans and L. acidophilus biofilms, except for 240 lactate production in the presence of raffinose. S. mutans and P. denticolens formed 241 more lactate than S. mutans or S. mutans and L. acidophilus biofilms in the presence of 242 243 glucose and sucrose. S. mutans and S. inopinata biofilms yielded more acetate and 244 lactate in the presence of all carbon sources. When raffinose was added to the culture 245 medium, S. mutans and S. inopinata biofilms produced 14 times more lactate and 48 times more acetate than S. mutans biofilms alone, even though this species participated 246 247 with 1.46% of the mixed biofilm (Tables 4 and 5).

248

249 Discussion

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The present study adds important information to the existing evidence in the literature. This is the first time that B. dentium, P. denticolens and S. inopinata are studied alone or in association with S. mutans, in either suspension or biofilms. Bacteria organized in biofilms are offered a higher antimicrobial resistance not only due its spatial organization – that impairs the penetration of antimicrobial substances – but also due to the low growth rate, phenotypical modifications and also because biofilms are extremely organized communities, in which interaction between cells confers an important resistance mechanism, as previously shown by Kara et al., 2006 [17].

Pilot studies showed that bifidobacteria are not able to form single species biofilms in the model used in the present study (data not shown). This is the reason why bifidobacteria single species biofilms were not evaluated in the present study. This is also a notable finding because it shows the importance of the interaction of bifidobacteria species with other oral microorganisms. More detailed study of bifidobacteria biofilms, including other analyses (i.e., confocal analyses), can generate important data on this interaction and should be conducted in the future.

The ability of bifidobacteria in suspension form to produce acids was already 266 demonstrated in previous studies. However, to the best of our knowledge this is the first 267 268 report on biofilms of bifidobacteria co-cultured with S. mutans. Haukioja et al., 2008 [18] showed that four different bifidobacteria strains were also able to promote a pH 269 drop below critical pH for enamel demineralisation (5.5) when different carbon sources 270 271 are used. Moynihan et al., 1998 [19] showed that B. dentium decreases culture medium 272 pH to values lower than enamel critical pH when exposed to glucose or lactose. Nakajo 273 et al., 2010 [20] also demonstrated the ability of bifidobacteria (B. dentium and Bifidobacterium longum) to decrease the pH culture below 5.0 at an initial pH of 5.0-274 275 7.0, indicating that these bacteria are able of creating an acidic environment in dental plaque and caries lesions. The acidogenic profile of bifidobacteria reaffirms their role in 276 the acidification of the oral environment, probably contributing in dental caries 277 278 development.

Carbon sources used in the present study were chosen based on their presence in the diet. Glucose, lactose and sucrose are either naturally present in fruits, vegetables or milk or added at high concentrations to baked products, snacks and sweets [21]. Raffinose is naturally present in beans, cabbage, brussels sprouts, broccoli, asparagus, and whole grains [21]. Xylitol is a natural sweetener that adds texture to foods and is not metabolized by most oral bacteria, including S. mutans [22]. The results of the present study support the evidence that bifidobacteria species present in carious lesions are able to metabolize all carbon sources included in the present study at different rates. Bifidobacteria demonstrated that they are able not only to produce significant amount of acids but also to accentuate biofilm acidogenicity in combination with S. mutans.

A possible explanation for the significant pH drop for the association between S. mutans and bifidobacteria is that S. mutans metabolizes carbon sources at a higher rate, producing acids more quickly than bifidobacteria and lactobacilli. Both lactobacilli and bifidobacteria prefer lower pH to produce acids, so acid production by S. mutans promotes a favorable environment to these species.

Bifidobacteria are able to metabolize raffinose to a higher extent than S. mutans and L. acidophilus, which reflected in a higher pH drop. This is an important finding because bifidobacteria do not require the consumption of snacks or sweets to produce acid, since raffinose is naturally present in healthy foods consumed on a daily basis. This can indicate the cariogenicity of bifidobacteria, which should be clinically evaluated.

Moreover, a synergistic effect between S. mutans and P. denticolens or S. 301 302 inopinata promoted a higher pH drop than these species alone. These results show that 303 the presence of both species in dental biofilm indicates a higher cariogenic potential 304 than if bifidobacteria are absent. This is of particular interest since some bifidobacteria 305 are used in probiotic foods. The use of probiotics on a daily basis is suggested to 306 modulate oral and intestinal microbiota. However, at the moment there are no clinical 307 trials that proved the beneficial use of bifidobacteria on caries prevention [23]. The 308 results of the present study suggest that the use of these species in probiotics may 309 increase pH drop and acid production in dental biofilm. The clinical outcome of these 310 findings should be further evaluated.

Our results for planktonic cultures demonstrated that S. inopinata was not able to ferment glucose and lactose efficiently, which is in disaccord with the literature [24]. Perhaps it simply ferments these two sugars slowly, in comparison to the rates of fermentation of raffinose and sucrose.

315 So, further studies on the metabolism of carbon sources should be performed not 316 only on the species investigated in this study, but also on other bifidobacteria such as Scardovia wiggsiae, which has recently been recognized as a member of the oral
microbiota [10,25]. Recently, higher prevalence of S. wiggsiae was found in caries
lesions than in controls [26] and this finding reinforce the need of deeper investigation
on other species.

Differences in pH drops and acid production observed in the control group might be related to inefficient carbohydrate depletion. So, the production of acids might be explained by the metabolism of residual endogenous polysaccharides.

In this study pH drop was evaluated by measuring the pH at baseline and after 3 h. Multiple measurements of pH over time may also generate interesting results and should be performed in future studies.

Overall, a higher amount of acetate was produced by bifidobacteria when 327 328 cultured planktonically, which is in agreement with findings reported by Crociani et al., 329 1996 [24]. This is also the first time that the fermentative profile of bifidobacteria in the 330 presence of lactose, raffinose and sucrose was studied in suspension and biofilms grown in the presence of S. mutans. Although acetate production is beneficial in the intestinal 331 environment, it may have detrimental effects in the oral cavity. Together with other oral 332 species, acetate production may contribute to environmental changes that shift healthy 333 334 oral microbiota to a more cariogenic one. More importantly, when cultured with S. mutans, bifidobacteria seem to contribute to a rise in lactate production, an important 335 feature in caries etiology. Also, other virulence factors, such as aciduricity, 336 337 antimicrobial resistance, and metabolic activity should be evaluated in the future.

Based on the findings of the present study, it is concluded that B. dentium, P. denticolens and S. inopinata are as acidogenic as S. mutans. Moreover, dual species biofilms of S. mutans and oral bifidobacteria produced a significantly greater pH drop than those produced by individual species.

342

343 Compliance with Ethical Standards

344

345 Conflict of Interest

346 Bruno Mello de Matos declares that he has no conflict of interest. Fernanda Lourenção

347 Brighenti declares that she has no conflict of interest. Thuy Do declares that she has no

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349	Yumi Koga Ito declares that she has no conflict of interest.
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355	
356	Ethical approval
357	All procedures performed in studies involving human participants were in accordance
358	with the ethical standards of the institutional and/or national research committee and
359	with the Declaration of Helsinki (1964) and its later amendments or comparable ethical
360	standards.
361	
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 inopinata. Int J Syst Evol Microbiol 61:25-29.
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- 456 26. Henne K, Rheinberg A, Melzer-Krick B, Conrads G (2015) Aciduric microbial taxa
 457 including Scardovia wiggsiae and Bifidobacterium spp. in caries and caries free
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459 Table 1: pH drop (Δ pH) (average ± sd) for single species and dual-species suspensions (n = 4)

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	glucose	lactose	raffinose	sucrose	control
S. mutans	2.06 ± 0.01	0.65 ± 0.01	1.06 ± 0.02	2.08 ± 0.01	0.53 ± 0.01
L. acidophilus	2.33 ± 0.03^a	0.64 ± 0.01	$0.94\pm0.01^{\texttt{a}}$	$1.88\pm0.03^{\rm a}$	0.45 ± 0.01
B. dentium	$2.18\pm0.02^{\text{a,b}}$	$1.12\pm0.11^{\text{a,b}}$	$2.19\pm0.02^{a,b}$	$2.21\pm0.01^{a,b}$	0.54 ± 0.02
P. denticolens	2.07 ± 0.01^{b}	$0.87\pm0.01^{\text{a},\text{b}}$	$1.55\pm0.04^{a,b}$	$1.56\pm0.01^{a,b}$	0.68 ± 0.04
S. inopinata	0.55 ± 0.01 ^{a,b}	0.57 ± 0.02	$2.01\pm0.06^{a,b}$	2.12 ± 0.06^{b}	0.49 ± 0.08
S. mutans + L. acidophilus	$2.21\pm0.01^{\text{a}}$	0.57 ± 0.01	$0.75\pm0.02^{\rm a}$	$2.28\pm0.04^{\rm a}$	0.34 ± 0.04
S. mutans + B. dentium	2.11 ± 0.11	$0.81\pm0.04^{\text{a,c}}$	$2.04\pm0.06^{\rm a,c}$	2.15 ± 0.10^{b}	0.48 ± 0.02
S. mutans + P. denticolens	$2.42\pm0.05^{a,c}$	$1.02 \pm 0.02^{a,c}$	$2.08\pm0.05^{\rm a,c}$	$2.44\pm0.04^{\rm a,c}$	0.93 ± 0.05
S. mutans + S. inopinata	$2.40\pm0.01^{\text{a,c}}$	$0.82\pm0.01^{\text{a,c}}$	$2.11\pm0.02^{\mathtt{a},c}$	$2.45\pm0.01^{\rm a,c}$	0.74 ± 0.01

461 Letters: significant differences within the same carbohydrate in relation to S. mutans (a), L. acidophilus (b), S. mutans + L. acidophilus (c); data in bold: no significant

differences in the same line in relation to control; ANOVA/Tukey's test, p<0.05

473 Table 2: pH drop (Δ pH) (average ± sd) for dual species biofilms (n = 4)

	glucose	lactose	raffinose	sucrose	control
S. mutans	1.66 ± 0.08	$\boldsymbol{0.86 \pm 0.05}$	1.28 ± 0.10	1.55 ± 0.05	0.76 ± 0.02
S. mutans + L. acidophilus	$1.97\pm0.03^{\rm a}$	0.93 ± 0.02	$1.72\pm0.06^{\text{a}}$	$2.02\pm0.01^{\text{a}}$	0.80 ± 0.01
S. mutans + B. dentium	1.56 ± 0.03^{b}	0.92 ± 0.01	1.30 ± 0.01^{b}	1.58 ± 0.08^{b}	0.79 ± 0.02
S. mutans + P. denticolens	$2.50\pm0.04^{a,b}$	$1.17 \pm 0.05^{a,b}$	$1.92\pm0.08^{\text{a,b}}$	$2.37\pm0.07^{a,b}$	1.05 ± 0.02
S. mutans + S. inopinata	$2.47\pm0.06^{a,b}$	$1.11 \pm 0.03^{a,b}$	$1.95\pm0.10^{\text{a,b}}$	$2.35\pm0.08^{a,b}$	0.99 ± 0.04

475 Letters: significant differences within the same carbohydrate in relation to S. mutans (a), S. mutans + L. acidophilus (b); data in bold: no significant differences in the same

476 line in relation to control; ANOVA/Tukey's test, p<0.05

477	Table 3:	Lactate and	acetate	concentration	(mM; averaged)	$ge \pm sd$)	in sus	pensions	(n = 8)	3)
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	gluo	cose	lact	tose	raffi	nose	suci	rose	con	trol
	lactate	acetate	lactate	acetate	lactate	acetate	lactate	acetate	lactate	acetate
S mutana	$3.56 \pm$	$0.36 \pm$	$0.00 \pm$	$0.02 \pm$	$0.39 \pm$	$0.32 \pm$	$3.73 \pm$	$0.32 \pm$	$0.00 \pm$	$0.01 \pm$
5. mutans	0.22	0.03	0.00	0.04	0.04	0.07	0.30	0.08	0.00	0.02
I a aidanhilua	$6.34 \pm$	$0.12 \pm$	$0.00 \pm$	$0.02 \pm$	$0.64 \pm$	$0.09 \pm$	0.19 ±	$0.02 \pm$	$0.00 \pm$	$0.00 \pm$
L. acidopinius	0.63ª	0.03ª	0.00	0.04	0.12ª	0.04ª	0.04 ^a	0.03 ^a	0.00	0.00
D. dontium	$0.91 \pm$	$2.74 \pm$	0.01 ±	$0.45 \pm$	$1.07 \pm$	$2.75 \pm$	$1.15 \pm$	$3.09 \pm$	$0.01 \pm$	$0.00 \pm$
D. dentium	$0.10^{a,b}$	0.13 ^{a,b}	0.02	$0.04^{a,b}$	$0.06^{a,b}$	0.18 ^{a,b}	$0.05^{a,b}$	0.16 ^{a,b}	0.03	0.00
D donticolons	$0.49 \pm$	$1.74 \pm$	$0.06 \pm$	$0.52 \pm$	$0.38 \pm$	1.11 ±	$0.43 \pm$	$1.16 \pm$	$0.19 \pm$	$0.57 \pm$
P. definicolefis	$0.05^{a,b}$	$0.12^{a,b}$	$0.04^{a,b}$	0.05 ^{a,b}	0.07^{b}	$0.11^{a,b}$	0.06^{a}	$0.08^{a,b}$	0.03	0.04
C inominata	$0.00 \pm$	$0.07 \pm$	$0.00 \pm$	$0.00 \pm$	$1.00 \pm$	$2.29 \pm$	$0.95 \pm$	$2.44 \pm$	$0.00 \pm$	$0.00 \pm$
S. mopinata	0.00 ^{a,b}	0.04 ^a	0.00	0.00	$0.10^{a,b}$	$0.17^{a,b}$	$0.08^{a,b}$	0.12 ^{a,b}	0.00	0.00
S. mutans +	$4.71 \pm$	$0.29 \pm$	$0.00 \pm$	$0.00 \pm$	$0.69 \pm$	$0.28 \pm$	$5.25 \pm$	$0.29 \pm$	$0.00 \pm$	$0.00 \pm$
L. acidophilus	0.36ª	0.04	0.00	0.00	0.08ª	0.05	0.17ª	0.05	0.00	0.00
S. mutans +	$3.18 \pm$	$1.82 \pm$	$0.00 \pm$	$0.03 \pm$	$1.32 \pm$	$2.47 \pm$	$4.21 \pm$	$2.94 \pm$	$0.00 \pm$	$0.00 \pm$
B. dentium	0.49 ^c	$0.10^{a,c}$	0.00	0.01	$0.08^{a,c}$	$0.22^{a,c}$	0.70°	$0.09^{a,c}$	0.00	0.00
S. mutans +	$5.66 \pm$	$1.97 \pm$	0.01 ±	$0.47 \pm$	$1.01 \pm$	$1.46 \pm$	$4.45 \pm$	$1.51 \pm$	$0.01 \pm$	$0.44 \pm$
P. denticolens	$0.29^{a,c}$	$0.11^{a,c}$	0.02	0.03 ^{a,c}	$0.09^{a,c}$	0.11 ^{a,c}	$0.22^{a,c}$	0.13 ^{a,c}	0.01	0.05
S. mutans +	$4.50 \pm$	$0.62 \pm$	$0.00 \pm$	$0.10 \pm$	$1.56 \pm$	$1.99 \pm$	$4.69 \pm$	$1.85 \pm$	$0.00 \pm$	$0.21 \pm$
S. inopinata	0.55ª	$0.04^{a,c}$	0.00	0.03 ^{a,c}	0.12 ^{a,c}	0.14 ^{a,c}	0.65 ^{a,c}	0.13 ^{a,c}	0.00	0.02

479 Letters: significant differences within the same carbohydrate in relation to S. mutans (a), L. acidophilus (b), S. mutans + L. acidophilus (c); data in bold: no significant differences in

480 the same line in relation to control; ANOVA/Tukey's test, p<0.05

481	Table 4: Lactate and acetat	e concentration (mM; avera	age \pm sd) in biofilms (n = 8)
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	gluc	cose	lact	cose	raffi	nose	suci	ose	con	trol
	lactate	acetate	lactate	acetate	lactate	acetate	lactate	acetate	lactate	acetate
S mutans	2.96 ±	0.00 ±	0.00 ±	0.00 ±	0.15 ±	0.03 ±	2.25 ±	0.00 ±	0.00 ±	0.00 ±
5. mutans	0.28	0.00	0.00	0.00	0.13	0.04	0.44	0.00	0.00	0.00
S. mutans +	2.15 ±	0.38 ±	$0.05 \pm$	0.22 ±	0.51 ±	$0.53 \pm$	3.23 ±	0.22 ±	$0.05 \pm$	$0.21 \pm$
L. acidophilus	0.49 ^a	0.12 ^a	0.05	0.08 ^a	0.19 ^a	0.22 ^a	1.18	0.09 ^a	0.06	0.20
S. mutans +	2.51 ±	0.09 ±	0.03 ±	0.06 ±	$0.40 \pm$	$0.23 \pm$	$2.28 \pm$	0.07 ±	$0.00 \pm$	$0.03 \pm$
B. dentium	0.19	0.06 ^b	0.07	0.08	0.09 ^a	0.05^{b}	0.59	0.06 ^b	0.00	0.06
S. mutans +	$9.50 \pm$	0.34 ±	0.07 ±	0.26 ±	$1.72 \pm$	$1.24 \pm$	$12.61 \pm$	0.34 ±	0.11 ±	$0.28 \pm$
P. denticolens	0.90 ^{a,b}	0.05 ^{a,b}	0.04	0.05 ^{a,b}	0.21 ^{a,b}	0.14 ^{a,b}	1.01 ^{a,b}	0.07 ^{a,b}	0.06	0.02
S. mutans +	9.23 ±	$0.55 \pm$	0.25 ±	0.45 ±	$2.16 \pm$	$1.46 \pm$	$11.79 \pm$	0.38 ±	$0.09 \pm$	$0.31 \pm$
S. inopinata	0.39 ^{a,b}	0.02 ^{a,b}	0.09 ^{a,b}	0.15 ^{a,b}	0.18 ^{a,b}	0.14 ^{a,b}	0.72 ^{a,b}	0.14 ^{a,b}	0.16	0.17

483 Letters: significant differences within the same carbohydrate in relation to S. mutans (a), S. mutans + L. acidophilus (b); data in bold: no significant differences in the same line in
 484 relation to control; ANOVA/Tukey's test, p<0.05

489 Table 5: Colony forming units (CFU/disc; average \pm sd) for dual species biofilms and specific species (lactobacilli and bifidobacteria) and 490 percentage of these species in relation to total count (%TM) (n=12)

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D: (1		Lactob	oacilli	Bifidobacteria		
BIOTIIM	l otal microorganism	CFU/disc	% TM	CFU/disc	%TM	
S. mutans	$2.77 \ge 10^7 \pm 1.44 \ge 10^7$	-	_	-	_	
S. mutans + L. acidophilus	7.09 x $10^7 \pm 2.83 \text{ x} 10^7$	$4.20 \ge 10^5 \pm 2.03 \ge 10^5$	0.59	-	-	
S. mutans + B. dentium	$5.15 \ge 10^6 \pm 2.18 \ge 10^6$	-	-	$4.66 \ge 10^6 \pm 2.19 \ge 10^6$	90.49	
S. mutans + P. denticolens	$2.83 \times 10^8 \pm 3.74 \times 10^7$	-	-	$3.84 \ge 10^6 \pm 1.67 \ge 10^6$	1.36	
S. mutans + S. inopinata	$2.52 \text{ x } 10^8 \pm 7.12 \text{ x } 10^7$	-	-	$3.68 \ge 10^6 \pm 1.50 \ge 10^6$	1.46	