

## Inhibition of *Streptococcus mutans* NS Adhesion to Glass with and without a Salivary Conditioning Film by Biosurfactant-Releasing *Streptococcus mitis* Strains

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Received 24 May 1999/Accepted 15 November 1999

**The release of biosurfactants by adhering microorganisms as a defense mechanism against other colonizing strains on the same substratum surface has been described previously for probiotic bacteria in the urogenital tract, the intestines, and the oropharynx but not for microorganisms in the oral cavity. Two *Streptococcus mitis* strains (BA and BMS) released maximal amounts of biosurfactants when they were grown in the presence of sucrose and were harvested in the early stationary phase. The *S. mitis* biosurfactants reduced the surface tensions of aqueous solutions to about 30 to 40 mJ m<sup>-2</sup>. Biochemical and physicochemical analyses revealed that the biosurfactants released were glycolipids. An acid-precipitated fraction was extremely surfactive and was identified as a rhamnolipidlike compound. In a parallel-plate flow chamber, the number of *Streptococcus mutans* NS cells adhering to glass with and without a salivary conditioning film in the presence of biosurfactant-releasing *S. mitis* BA and BMS (surface coverage, 1 to 4%) was significantly reduced compared with the number of *S. mutans* NS cells adhering to glass in the absence of *S. mitis*. *S. mutans* NS adhesion in the presence of non-biosurfactant-releasing *S. mitis* BA and BMS was not reduced at all. In addition, preadsorption of isolated *S. mitis* biosurfactants to glass drastically reduced the adhesion of *S. mutans* NS cells and the strength of their bonds to glass, as shown by the increased percentage of *S. mutans* NS cells detached by the passage of air bubbles through the flow chamber. Preadsorption of the acid-precipitated fraction inhibited *S. mutans* adhesion up to 80% in a dose-responsive manner. These observations indicate that *S. mitis* plays a protective role in the oral cavity and protects against colonization of saliva-coated surfaces by cariogenic *S. mutans*.**

Dental plaque is a complex, multispecies biofilm that forms on oral surfaces in which bacteria are embedded in bacterial and salivary polymers. Many species and strains have been identified in dental plaque; some of these organisms (29, 32) are now recognized as early colonizers, and others (16, 34) are considered cariogenic organisms or periodontopathogens (36). However, no real ecological role has been defined for most strains and species in dental plaque, and it is possible that these organisms should be considered part of the normal, indigenous, healthy microflora in the oral cavity.

Similarly, normal healthy microfloras have been identified for the gastrointestinal tract (31), the urogenital tract (17, 24), the skin (15), and the eyes (18). In a healthy host, a normal microflora effectively competes with invading pathogens. The mechanisms employed by the healthy microflora to interfere with the adhesion of invading pathogens include competitive exclusion (26) and displacement (20), production of antibacterial compounds, such as lactic acid, hydrogen peroxide, bacteriocins, and bacteriocin-like substances, by lactobacilli (13, 19), coaggregation (7, 25), and release of biosurfactants (8).

Recently, it has been demonstrated that biosurfactants released by lactobacilli can be adsorbed to catheter materials in order to discourage uropathogen adhesion (37). Microorganisms in certain dairy products also release biosurfactants that inhibit adhesion of yeasts, most notably to voice prostheses in the oropharynx (1). Several years ago, it was pointed out that *Streptococcus mitis* BMS released substances, which were later recognized as biosurfactants (33), that discourage adhesion of

*Streptococcus mutans* (23). This observation has never been followed up on, but such a process is of considerable interest as a mechanism that could be used to prevent dental caries, since *S. mutans* is an important etiological agent of coronal (16, 34) and root surface caries (30).

The aim of this study was twofold: (i) to determine the surface tensions of biosurfactant solutions and their chemical characteristics when they are released by two oral *S. mitis* strains grown on different carbohydrate sources and (ii) to determine the effect of biosurfactant-releasing *S. mitis* cells adhering to an artificial (glass) substratum with and without a salivary conditioning film on the subsequent adhesion of a *S. mutans* strain. In addition, surface properties of *S. mitis* cells before and after biosurfactants were released were measured in order to rule out the possibility that the compounds released were cell surface compounds whose release affected the adhesive properties of the cell surface.

### MATERIALS AND METHODS

**Microorganisms.** *S. mutans* NS and *S. mitis* BA and BMS were originally isolated from the human oral cavity and were stored in Todd-Hewitt broth (THB) (Oxoid, Basingstoke, England) supplemented with 0.5% sucrose and 7% (vol/vol) dimethyl sulfoxide at -60°C. Streptococci from the frozen stock preparations were streaked every 2 weeks onto blood agar plates and incubated at 37°C. After 2 days the plates were stored at 5°C.

For adhesion assays and surface characterization studies the bacteria from a blood agar plate were grown overnight at 37°C in 10 ml of THB supplemented with 0.5% sucrose. The resulting culture was used to inoculate a second culture, which was grown for 16 h and then harvested by centrifugation at 4,000 × g, washed twice with adhesion buffer (2 mM potassium phosphate, 50 mM potassium chloride, 1 mM calcium chloride; pH 6.8), and resuspended in adhesion buffer.

To break bacterial chains, a bacterial suspension was sonicated five times for 10 s at 30 W (model 375 Vibra Cell sonicator; Sonics and Materials Inc., Danbury, Conn.). Sonication was performed intermittently while the preparation was cooling in an ice-water bath. Finally, the cells were suspended in adhesion

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buffer. The cell concentration was fixed at  $3 \times 10^8$  cells ml of buffer<sup>-1</sup> by using a Bürker-Türk counting chamber.

**Biosurfactant release.** To release biosurfactant from *S. mitis* BA and BMS, subcultures (10 ml) from blood agar plates were prepared by inoculating THB supplemented with 0.5% (wt/vol) glucose, 0.5% (wt/vol) glycerol, 0.5% (wt/vol) galactose, or 0.5% (wt/vol) sucrose and incubating the preparations overnight at 37°C.

An overnight subculture was used to inoculate 1,400 ml of a second culture. Cells were harvested in the mid-exponential, early-stationary, and stationary phases by centrifugation at  $4,000 \times g$ , washed twice in adhesion buffer, and resuspended in 200 ml of water. Crude biosurfactant was produced by gently stirring the suspension for 2 h at room temperature. Subsequently, the microorganisms and the biosurfactants released were separated by centrifugation at  $10,000 \times g$ . To ensure that all of the cell remnants were removed, the supernatant was centrifuged twice at  $10,000 \times g$ . After the final centrifugation, both the cellular pellet and the crude biosurfactant were freeze-dried and weighed, and the crude biosurfactant was stored at  $-20^\circ\text{C}$  until it was used.

In a separate experiment, the freeze-dried crude biosurfactant of *S. mitis* BMS was resuspended in water and subsequently acid precipitated with concentrated HCl at pH 2.0. After the supernatant was decanted, the precipitate was washed twice with acidic water (pH 2) and collected by centrifugation at  $4,000 \times g$ . After the acid precipitate was redissolved, it was freeze-dried. The supernatant was adjusted to pH 7 with KOH and also freeze-dried.

**Surface tension measurements.** Axisymmetric drop shape analysis by profile (ADSA-P) was performed as described by Noordmans and Busscher (22) in order to determine the surface tensions of biosurfactant solutions. Briefly, ADSA-P involves digitizing the circumference of a liquid droplet on a solid surface. The circumference of the droplet is fitted to the Laplace equation of capillarity (27), which yields the surface tension of the biosurfactant solution. Droplets containing biosurfactant dissolved in water (volume, approximately 100  $\mu\text{l}$ ) were placed on a clean piece of fluoroethylenepropylene (Teflon). Measurements for one solution droplet were obtained after 2 h in order to allow equilibration of the interface in an enclosed chamber at room temperature. One liquid profile was recorded twice with a minimal time interval ( $<0.5$  s) between measurements, and the ADSA-P surface tensions were averaged. This procedure was performed in duplicate with separate liquid droplets.

ADSA-P surface tension measurements were obtained for crude biosurfactant solutions in water as a function of biosurfactant concentration, as well as for the freeze-dried acid-precipitated fraction and supernatant of crude *S. mitis* BMS biosurfactant.

**Biochemical assay.** The protein content of the crude biosurfactant was determined by the Bio-Rad protein assay; bovine albumin was used as the standard.

**XPS.** For X-ray photoelectron spectroscopy (XPS), 100- $\mu\text{l}$  droplets of crude stationary-phase biosurfactants and the acid-precipitated fraction dissolved in water (approximately 10 mg ml<sup>-1</sup>) were placed on gold-coated glass slides (1 by 1 cm). After air drying, the glass slides were inserted into the chamber of a spectrometer (Surface Science Instruments, S-probe, Mountain View, Calif.). The residual pressure in the spectrometer during operation was approximately  $10^{-9}$  Pa. A magnesium anode was used to produce X-rays (10 kV, 22 mA) with a spot size of 250 by 1,000  $\mu\text{m}$ . After scans of the overall spectrum in the binding energy range from 1 to 1,200 eV at low resolution (150-eV pass energy) were obtained, peaks over a 20-eV binding energy range were recorded at high resolution (50-eV pass energy) in the following order: C<sub>1s</sub> (four scans), O<sub>1s</sub> (four scans), N<sub>1s</sub> (eight scans), P<sub>2p</sub> (eight scans), and C<sub>1s</sub> again in order to account for contamination or deterioration under X rays of the samples.

The carbon peak was split by a least-squares fitting program into four Gaussian components at 284.8, 286.2, 287.8, and 289.2 eV by imposing a constant full width at a half-maximum of 1.35 eV, and these components were thought to be representative of carbon involved in C—C, C—O and C—N, C=O and O—C—O, and O=C—OH bonds, respectively. The oxygen peak was split into two components at 531.0 and 532.4 eV by imposing a constant full width at a half-maximum of 1.70 eV, and these components were thought to be representative of oxygen involved in O=C and C—O bonds, respectively.

**Cell surface properties.** The bacterial cell surface hydrophobicity of the *S. mitis* strains was assessed by measuring water contact angles on bacterial lawns (35) on membrane filters (pore diameter, 0.45  $\mu\text{m}$ ). Wet filters with deposited organisms were fixed on sample holder plates with double-sided sticky tape and air dried. Water contact angles were measured after 20 min by using image analysis techniques at 25°C and sessile droplets of water. At least three different filters containing samples from separate cultures were prepared.

The zeta potentials of the *S. mitis* strains were determined in adhesion buffer from the speed of suspended bacteria in a 150-V applied electric field by using the Helmholtz-Smoluchowski equation (12). The instrument used, a model 501 Lazer Zee meter (PenKem, Bedford Hills, N.Y.), was equipped with an image analysis option for tracking and zeta sizing (38).

**Saliva.** Human whole saliva was collected from 10 healthy volunteers of both sexes in ice-chilled cups. Saliva production by the volunteers was stimulated by chewing Parafilm (3M Company, Minneapolis, Minn.). After the saliva was pooled and centrifuged at  $10,000 \times g$  for 5 min at 10°C, phenylmethylsulfonyl fluoride (0.2 M; Merck, Darmstadt, Germany) was added to a final concentration of 1 mM as a protease inhibitor. The solution was centrifuged again, dialyzed overnight at 4°C against water, and freeze-dried for storage. A 1.5-mg ml<sup>-1</sup>

TABLE 1. Cell dry weights and amounts of biosurfactant released in different growth phases by *S. mitis* strains grown in media supplemented with different carbohydrate sources, expressed as averages of data from three experiments performed with separately cultured bacteria<sup>a</sup>

Strain	Growth phase	Carbohydrate source	Cell dry wt (g liter of medium <sup>-1</sup> )	Biosurfactant yield (mg g <sup>-1</sup> )
<i>S. mitis</i> BA	Stationary	Glucose	0.50	36
		Glycerol	0.25	32
		Galactose	0.38	26
		Sucrose	0.58	31
		Sucrose	0.50	52
<i>S. mitis</i> BMS	Early stationary	Sucrose	0.08	125
		Sucrose	0.08	125
<i>S. mitis</i> BMS	Stationary	Glucose	0.43	28
		Glycerol	0.18	56
		Galactose	0.35	37
	Early stationary	Sucrose	0.40	30
		Sucrose	0.38	42
	Mid-exponential	Sucrose	0.10	70
		Sucrose	0.10	70

<sup>a</sup> In all cases the standard deviation was less than 15%.

solution of freeze-dried stock in adhesion buffer (see above) was designated reconstituted human whole saliva.

**Adhesion experiments.** The flow chamber (length, 7.6 cm; width, 3.8 cm; height, 0.06 cm) and image analysis system used have been described in detail previously (28). Images were obtained from the bottom glass plate (5.8 by 3.8 cm) of the parallel-plate flow chamber. The top plate of the chamber was also made of glass.

Prior to each experiment, all tubes and the flow chamber were filled with adhesion buffer, and care was taken to remove air bubbles from the system. Flasks containing microbial suspensions, buffer, and saliva when appropriate were placed at the same height with respect to the chamber, so that immediately after the flow was switched, all of the fluids would circulate through the chamber under the influence of hydrostatic pressure at a shear rate of  $20 \text{ s}^{-1}$  ( $0.05 \text{ ml s}^{-1}$ ; well within the limits of laminar flow), which represented a shear rate between the shear rates exerted by stimulated and unstimulated salivary flow at the level of the tooth surface (4). First, the flow was switched to saliva (when appropriate) for 1.5 h in order to create a salivary conditioning film, and then the flow was switched for 15 min to buffer to remove all remnants of saliva from the tubing and the flow chamber. After this, the flow was switched (when appropriate) to an *S. mitis* BMS or BA suspension until the desired surface coverage by adhering *S. mitis* BMS or BA cells (between 1 and 4%) was attained, as measured in real time with the image analysis system. In the experiments carried out to determine the effects of preadsorbed biosurfactants, biosurfactants in a 20-mg ml<sup>-1</sup> solution were adsorbed overnight to the glass plate. Between flow steps, buffer was run through the system to remove unbound material from the tubes and chamber. Finally, a *S. mutans* suspension was circulated through the system for 4 h.

The initial increase in the number of adhering *S. mutans* NS cells with time was expressed by a so-called initial deposition rate; this rate was the number of microorganisms that adhered initially per unit of time and area. The number of bacteria adhering after 4 h was considered an estimate of microbial adhesion at a more advanced stage of the adhesion process.

Finally, after 4 h, air bubbles were passed through the chamber in order to obtain an indication of the adhesive forces (14). The passage of an air-liquid interface (i.e., an air bubble) over adhering micron-sized particles is accompanied by a detachment force of about  $10^{-7}$  N per adhering microorganism. After the passage of each air bubble, the mean percentage of bacteria detached by the high removal force was determined at five different spots on the substratum surface with respect to the mean number of organisms adhering at five spots prior to the introduction of the first air bubble.

All adhesion experiments were performed in triplicate with separately cultured organisms at room temperature.

## RESULTS

Table 1 shows the amounts of biosurfactant released by *S. mitis* BA and BMS grown on different carbohydrates in various growth phases. The amount of biosurfactant released per gram of dry cell weight was largest for bacteria in the mid-exponential growth phase and decreased as the organisms entered the stationary growth phase. Adding glycerol to the medium increased the biosurfactant yield of *S. mitis* BMS in the stationary phase compared to the other carbohydrates used. *S. mitis*

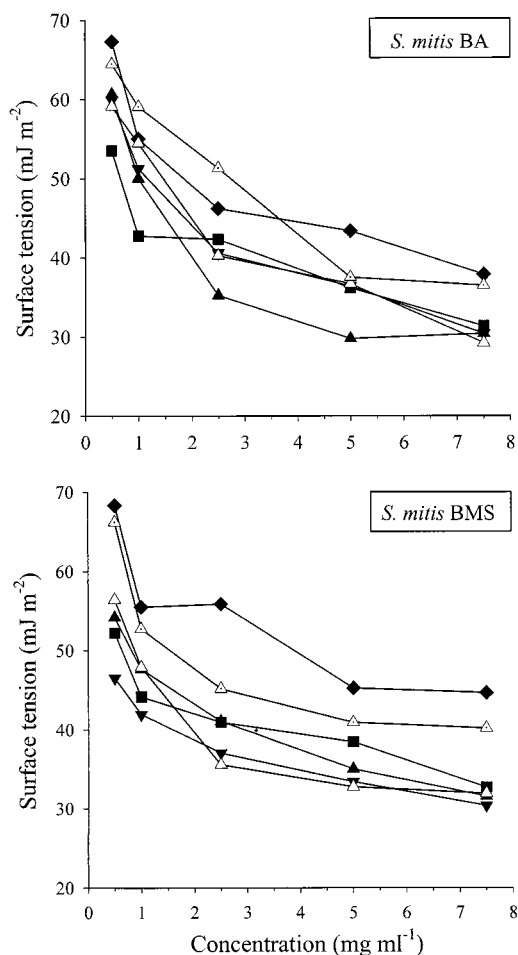


FIG. 1. Plots of surface tensions of biosurfactant solutions in water versus concentrations of freeze-dried biosurfactants released by *S. mitis* BA and BMS grown on medium supplemented with glucose (◆), glycerol (■), galactose (▼), or sucrose (▲, △, and ▽) and harvested in the mid-exponential phase (△), early-stationary phase (▽), and stationary phase (◆, ■, ▼, and ▲). The results are averages of data obtained from three separately grown cultures with standard deviations of less than 10%.

BA in the stationary growth phase released similar amounts of biosurfactants irrespective of the carbohydrate present.

Figure 1 shows that the *S. mitis* biosurfactants could reduce the surface tension of an aqueous solution to around 30 to 40 mJ m<sup>-2</sup> irrespective of the growth phase or the carbohydrate present. When glucose was used as the carbohydrate, the surface activity of the compounds released was less than the surface activities after growth on the other carbohydrates used.

The protein contents of the biosurfactants released by both *S. mitis* BA and BMS were extremely low (less than 1%). Also, XPS physicochemical analyses (Table 2) revealed that the biosurfactants contained little nitrogen (the N/C ratios were between 0.101 and 0.172) compared with protein reference compounds (the N/C ratio is 0.270 for the average protein). The nitrogen contents of the biosurfactants were too high, however, for association of the biosurfactants with lipoteichoic acid, which was also ruled out by relatively low oxygen and phosphorus contents. The absence of proteins and lipoteichoic acids in the biosurfactants was confirmed by the low fractions of carbon doubly bound to oxygen (C=O). C—(C,H) functionalities were abundant in the biosurfactants released by the *S. mitis* strains, as they are in rhamnolipids.

Table 3 shows the adhesion of *S. mutans* NS in the presence of adhering, biosurfactant-releasing *S. mitis* strains grown on sucrose. As Table 3 shows, when the surface coverage by *S. mitis* BMS on bare glass was up to 4%, the initial deposition rate of *S. mutans* NS was reduced from 1,043 to 394 cm<sup>-2</sup> s<sup>-1</sup>. Also, the number of *S. mutans* cells adhering after 4 h decreased from 130 × 10<sup>5</sup> to 50 × 10<sup>5</sup> cells cm<sup>-2</sup>. The effects of biosurfactant-releasing *S. mitis* BA on *S. mutans* NS adhesion were far less obvious than the effects of *S. mitis* BMS. Whereas surface coverage by *S. mitis* BA of up to 2% resulted in similar or sometimes even greater reductions in *S. mutans* adhesion than the reductions observed in the presence of *S. mitis* BMS, the effect of *S. mitis* BA on *S. mutans* adhesion completely disappeared when the surface coverage was 4%. Finally, *S. mitis* BA and BMS biosurfactants decreased the strength of *S. mutans* NS bonds to glass, as the percentage of adhering *S. mutans* cells that were detached by air bubble passage increased as the surface coverage by the *S. mitis* strains increased.

In order to determine whether the decreases in *S. mutans* NS

TABLE 2. Chemical composition data (as determined by XPS) for *S. mitis* BMS and BA biosurfactants released in the stationary growth phase and for reference compounds

Crude biosurfactant or compound	Elemental concn ratios			Fractions of carbon and oxygen involved in various chemical functions					
	N/C	O/C	P/C	C—(C,H)	C—(O,N)	C=O	O=C—OH	(O=C)/C	(OH)/C
Biosurfactant released by:									
<i>S. mitis</i> BA	0.172	0.407	0.019	0.58	0.24	0.18	0	0.342	0.065
<i>S. mitis</i> BMS	0.101	0.426	0.016	0.64	0.20	0.13	0.04	0.268	0.158
Acid-precipitated <i>S. mitis</i> BMS biosurfactant	0.071	0.366	0.016	0.61	0.24	0.11	0.05	0.121	0.244
Reference compounds									
Protein <sup>a</sup>	0.270	0.32	0	0.41	0.32	0.28 <sup>f</sup>		0.275	0.045
Glycosidic residue <sup>b</sup>	0	0.83	0	0	0.83	0.17 <sup>f</sup>		0	0.833
Rhamnolipid R1 <sup>c</sup>	0	0.406	0	0.56	0.31	0.06	0.06	0.061	0.345
Rhamnolipid R2 <sup>c</sup>	0	0.346	0	0.65	0.23	0.04	0.08	0.076	0.269
Phospholipid <sup>d</sup>	0.009	0.21	0.034	0.82	0.13	0.05	0.01	0.077	0.132
LTA <sup>e</sup>	0.031	0.63	0.066	0.41	0.44	0.44	0.10	0.151	0.478

<sup>a</sup> Average protein, calculated based on a collection of bacterial, fungal, and mammalian proteins (21).

<sup>b</sup> Glycosidic residue C<sub>6</sub>H<sub>10</sub>O<sub>5</sub> (21).

<sup>c</sup> Calculated for rhamnolipids synthesized by *Pseudomonas aeruginosa*. R1 contains two rhamnose units, and R2 contains one rhamnose unit (6, 11).

<sup>d</sup> L- $\alpha$ -Phosphatidyl-DL-glycerol dimyristoyl.

<sup>e</sup> LTA, Lipoteichoic acid from *S. mutans* (Sigma).

<sup>f</sup> C=O and O=C—OH together.

TABLE 3. Adhesion of *S. mutans* NS to glass in competition with adhering biosurfactant-releasing and nonreleasing *S. mitis* BA and BMS and to glass with preadsorbed stationary-phase *S. mitis* BA and BMS biosurfactants

Prepn	Surface coverage	Initial deposition rate (cm <sup>-2</sup> s <sup>-1</sup> )	No. of bacteria adhering after 4 h (10 <sup>5</sup> cells cm <sup>-2</sup> )	Detachment due to air bubbles (%) <sup>a</sup>		
				First bubble	Second bubble	Third bubble
Glass without salivary conditioning film	None	1,043 ± 160 <sup>b</sup>	130 ± 18	14 ± 16	20 ± 11	32 ± 9
	1% <i>S. mitis</i> BA	798 ± 148	111 ± 5	0 ± 10	5 ± 16	0 ± 16
	2% <i>S. mitis</i> BA	577 ± 331	73 ± 43	0 ± 19	7 ± 12	19 ± 20
	4% <i>S. mitis</i> BA	1,096 ± 127	100 ± 4	5 ± 12	27 ± 13	47 ± 19
	<i>S. mitis</i> BA biosurfactant alone	48 ± 49	4 ± 3	47 ± 8	74 ± 9	67 ± 16
	4% <i>S. mitis</i> BA <sup>-c</sup>	977 ± 279	111 ± 26	1 ± 6	8 ± 18	11 ± 9
	1% <i>S. mitis</i> BMS	774 ± 65	100 ± 15	15 ± 13	16 ± 20	20 ± 15
	2% <i>S. mitis</i> BMS	895 ± 293	92 ± 21	42 ± 19	17 ± 17	39 ± 50
	4% <i>S. mitis</i> BMS	394 ± 83	50 ± 11	45 ± 21	35 ± 15	43 ± 9
	<i>S. mitis</i> BMS biosurfactant alone	106 ± 37	9 ± 4	59 ± 11	57 ± 21	68 ± 16
	4% <i>S. mitis</i> BMS <sup>-</sup>	789 ± 168	113 ± 28	12 ± 25	0 ± 11	0 ± 15
	Glass with salivary conditioning film	None	965 ± 100	83 ± 9	72 ± 20	70 ± 22
1% <i>S. mitis</i> BA		495 ± 331	37 ± 11	72 ± 19	77 ± 15	77 ± 21
1% <i>S. mitis</i> BMS		657 ± 277	49 ± 20	72 ± 15	92 ± 5	91 ± 7
2% <i>S. mitis</i> BMS		573 ± 262	43 ± 19	70 ± 22	78 ± 9	82 ± 12
4% <i>S. mitis</i> BMS		349 ± 35	30 ± 8	60 ± 16	67 ± 8	61 ± 18

<sup>a</sup> Total percentages of *S. mutans* NS cells that were detached by the passage of three consecutive air bubbles through the flow chamber.

<sup>b</sup> The data are means ± standard deviations based on data from three experiments performed with separately cultured cells.

<sup>c</sup> The minus sign denotes nonreleasing cells.

adhesion in the presence of adhering *S. mitis* strains were due to simple geometrical effects or to the biosurfactants released by the adhering *S. mitis* strains, we performed experiments with two non-biosurfactant-releasing preparations, *S. mitis* BA<sup>-</sup> and BMS<sup>-</sup>. Nonreleasing bacteria were prepared by allowing the *S. mitis* strains to release their biosurfactants overnight; after this they were used in adhesion experiments with *S. mutans* NS. Using ADSA-P surface tension measurements, we determined that the BA<sup>-</sup> and BMS<sup>-</sup> bacteria had lost the ability to reduce the surface tension of an aqueous suspension ( $\Delta\gamma = 0$  and 2 mJ m<sup>-2</sup>, respectively) and thus their ability to release biosurfactants. Furthermore, we observed no significant changes in the cell surface hydrophobicities and zeta potentials of the organisms that might affect *S. mutans* adhesion. The water contact angle on *S. mitis* BMS was 99° before and after the biosurfactant was released, while the water contact angle on *S. mitis* BA was 104°. Similarly, the zeta potential of *S. mitis* BMS in adhesion buffer was not altered by biosurfactant release; it remained -9 mV. For *S. mitis* BA, there was an insignificant change in the bacterial zeta potential from -8 to -5 mV that occurred when the biosurfactant was released.

Table 3 shows that there was not a significant difference between adhesion of *S. mutans* NS in the presence of adhering, nonreleasing *S. mitis* BA<sup>-</sup> and BMS<sup>-</sup> (surface coverage, 4%) and adhesion to bare glass. Table 3 also shows that preadsorption of crude biosurfactants released by *S. mitis* BA, as well as by *S. mitis* BMS, significantly reduced the adhesion of *S. mutans* to glass. The initial deposition rates and the numbers of *S. mutans* cells adhering after 4 h to biosurfactant-coated glass were almost 10-fold lower in the presence of *S. mitis* BMS biosurfactant and more than 20-fold lower in the presence of *S. mitis* BA biosurfactant. Also, detachment of adhering *S. mutans* cells by air bubble passage increased drastically in the presence of preadsorbed crude biosurfactants.

The acid-precipitated fraction of *S. mitis* BMS biosurfactant was extremely surface active compared with the crude biosurfactant, and an aqueous solution of the acid precipitate containing only 1 mg ml<sup>-1</sup> had a surface tension of 35 mJ m<sup>-2</sup> (compare with Fig. 1) while an aqueous solution of the freeze-dried supernatant (the supernatant left after acid precipitation) hardly decreased the surface tension of water in the same

concentration range. Moreover, substratum coverage by adsorption of the acid-precipitated fraction from an aqueous solution (between 0.1 and 1 mg ml<sup>-1</sup>) resulted in up to 80% inhibition of *S. mutans* NS adhesion in a dose-responsive manner. XPS analysis of the acid-precipitated fraction revealed a reduced nitrogen content compared to the crude biosurfactant (Table 2), while the O/C elemental concentration ratio was between the ratios for rhamnolipids R1 and R2.

Table 3 also shows the adhesion of *S. mutans* NS to glass with a salivary conditioning film in the presence of stationary-phase biosurfactant-releasing *S. mitis* BA and BMS grown on sucrose. This table shows that biosurfactant-releasing *S. mitis* strains interfered not only with *S. mutans* adhesion to glass but also with *S. mutans* adhesion to salivary conditioning films.

## DISCUSSION

In this study we found that release of biosurfactants by adhering *S. mitis* strains interferes with adhesion of a cariogenic *S. mutans* strain. A chemical characterization study suggested that glycolipids were present in the crude biosurfactant, while further purification revealed that the active component was rhamnolipidlike. It was found previously that *S. mitis* ATCC 9811 released extracellular substances that were surface active (10, 33). However, these extracellular substances were identified as an exohemagglutinin, a lectinlike substance that is responsible for interaction with saliva (10).

The most commonly isolated biosurfactants are glycolipids (e.g., rhamnolipids produced by *Pseudomonas aeruginosa* [9]) and lipopeptides (e.g., surfactin released by *Bacillus subtilis* [3]). The yields of both of these types of biosurfactants are relatively high (approximately 2.5 g liter of medium<sup>-1</sup>), and these biosurfactants reduce the surface tension of spent culture supernatant to less than 30 mJ m<sup>-2</sup>. *Streptococcus thermophilus* B (2) and *Lactobacillus* species (37) are also biosurfactant-releasing strains. The biosurfactants of these organisms decrease the surface tension of water to around 37 mJ m<sup>-2</sup>, but the amounts released per liter of culture medium were orders of magnitude smaller than the amounts of the rhamnolipids and lipopeptides released (approximately 100 mg liter<sup>-1</sup> for *Lactobacillus* species and 20 mg liter<sup>-1</sup> for *S. thermophilus* B).

A simple calculation revealed that small amounts of biosurfactants may have substantial effects on adhesion to substratum surfaces. If it is assumed that the biosurfactant is a small molecule with a molecular weight of about 1,000, it can be estimated based on a biosurfactant yield of around  $10^{-8}$  mg per cell that a substratum surface coverage value of around 8% for biosurfactant-releasing *S. mitis* BA or BMS cells results in 100% coverage of the surface by biosurfactants (2). Indeed, the adhesion experiments performed with *S. mitis* BA and BMS grown on medium supplemented with sucrose showed that the presence of biosurfactant-releasing *S. mitis* BA and BMS cells on glass and on glass with a salivary conditioning film effectively reduced the adhesion of *S. mutans* NS even with the influence of geometrical effects (i.e., physical collisions between suspended and adhering organisms which resulted in increased deposition) (Table 3). Moreover, preadsorption of the biosurfactants to glass also drastically reduced *S. mutans* NS adhesion.

With regard to the oral cavity, the increased detachment of adhering *S. mutans* cells in the presence of biosurfactants after air bubble passage is very important, as air-liquid interfaces frequently pass over the enamel surface and the adhering microorganisms during eating and swallowing (i.e., adhering microorganisms are exposed to high detachment forces). Our results indicate that even if *S. mutans* NS adheres to bare surfaces or to a salivary conditioning film, biosurfactants effectively stimulate detachment of this organism by the dynamic shear forces that occur in the oral cavity.

In conclusion, adhering *S. mitis* BA and BMS cells decrease the adhesion of *S. mutans* NS to glass and to glass with a salivary conditioning film through the release of rhamnolipidlike biosurfactants. Thus, we propose that *S. mitis* plays an ecological role in the oral cavity.

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