1 2 3 4 5 6 7 8	Mucosal environment induces phage susceptibility in <i>Streptococcus mutans</i> Lotta-Riina Sundberg <sup>1*</sup> , Noora Rantanen <sup>1</sup> , Gabriel Magno de Freitas Almeida <sup>2</sup>
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14 15 16 17	Key words: bacteriophage, biofilm, mucin, <i>Streptococcus mutans</i> , phage, phage therapy
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24 25 26 27 28 29	<b>Authorship contribution statement</b> Conceptualization: GMFA, L-RS. Methodology: GMFA, NR, L-RS. Data curation: GMFA, NR, L-RS. Writing- Original draft preparation: LRS. Writing – review and editing: : GMFA, NR, L-RS. Visualization: GMFA, L-RS. Supervision, project administration, and funding acquisition: L-RS.

#### **Abstract**

Pathogenic bacteria are attracted towards mucosa, as it is their way of entry into the body. Yet, we know surprisingly little about the phage-bacterium interactions in the mucosal environment. Here, we explored the effect of the mucosal environment on growth characteristics and phage-bacterium interactions in *Streptococcus mutans*, causative agent of dental caries. We found that while mucin supplementation increased bacterial growth and survival, it decreased *S. mutans* biofilm formation. More importantly, presence of mucin had a significant effect on *S. mutans* phage susceptibility. In two experiments done in BHI, phage M102 replication was detected only with 0.2% mucin supplementation. In 0.1xTSB 0.5% mucin supplementation led to 4-log increase in phage titers compared to control. These results suggest that the mucosal environment can have a major role for growth, phage sensitivity and phage resistance of *S. mutans*, and underline the importance of understanding the effect of mucosal environment on phage-bacterium interactions.

#### Introduction

Many pathogenic bacteria use mucosal layers to invade metazoan hosts. Indeed, the mucosa is an important chemotactic signal for many pathogenic bacteria <sup>1,2</sup>, directing the bacteria towards the metazoan host. Mucosal cues cause expressional changes in virulence-related genes of many bacterial species <sup>3</sup>. For example, interaction with mucus and mucin glycoproteins impacts bacterial phenotypic characteristics and increases virulence of many pathogenic species <sup>4–6</sup>, including *Acinetobacter baumannii* <sup>7</sup> and *Streptococcus mutans* <sup>8</sup>. Furthermore, some bacteria, such as pathogens and many intestinal anaerobes, can use mucosal components directly as a nutrient source <sup>5</sup>.

The mucosal surface consists of secreted mucin glycoproteins, that oligomerize to form a complex viscous milieu allowing retention of antimicrobial molecules (e.g. immunoglobulins A and G, lectins, antimicrobial peptides) and also phages. Over 20 families of mucin glycoproteins exist, and they are expressed at a tissue-specific manner <sup>1,9</sup>. Mucin secretion is primarily constitutive, leading to a dynamic gradient of mucins - high close to the epithelial cell surface but lower in the outer surface of the mucosa.

The last 10 years of research have clearly shown that phages are major partners in the mucosal microbiomes <sup>10,11</sup>, thus affecting both the invading and local bacterial populations. Understanding the ecology of pathogens and their phages in the mucosa is central for our health, and for developing of phage therapy approaches against antibiotic resistant infections. Yet, we know surprisingly little about the phage-bacterium interactions in the mucosal environment. This has led to fundamental gaps and biases in our knowledge regarding the antagonistic interactions between phages and bacteria as they happen in the mucosal milieu.

The oral environment is an important habitat for phage-bacterium interactions<sup>12,13</sup>. The salivary microbial community has been shown to host 10<sup>8</sup> virus-like particles per ml (mostly dsDNA phages) <sup>14</sup> whereas in the dental plaque the density is higher, 10<sup>10</sup> virus-like particles per gram <sup>15</sup>. Since previous research suggests mucosa may influence the virulence of bacteria <sup>4-6</sup> and phages <sup>6,16,17</sup>, and favours phage resistance mechanisms that maintain bacterial virulence (e.g. CRISPR-Cas) <sup>18</sup>, it is important to describe how the mucosal environment shapes these interactions in *Streptococcus mutans*, causative agent of dental caries<sup>18</sup>

Here, we characterized the effect of mucosal environment on growth features and phage-bacterium interactions in *S. mutans*. In its natural habitat, *S. mutans* forms biofilm and is in continuous contact with oral mucosal surfaces. It has been previously demonstrated, that several salivary components inhibit *S. mutans* biofilm formation, but the presence of sucrose

can reverse this effect <sup>19</sup>. Yet, so far there has been no direct demonstration of the impact of mucins on the interactions between *S. mutans* and its phages This aspect is central for understanding the phage-bacterium interactions in the mucosa, and for developing successful phage therapy approaches against caries. Here, we characterized the impact of mucosal environment on *S. mutans* growth characteristics in physiologically relevant mucin concentrations <sup>20,21</sup>. We found that while mucin supplementation decreases *S. mutans* biofilm formation, it increases bacterial survival in starvation. More importantly, mucin has a central role for *S. mutans* susceptibility to phage infections, and, consequently, phage resistance.

### Materials and methods

### Host and phage strains

Streptococcus mutans OMZ381 and its phage M102 were obtained from Félix d'Hérelle Reference Center for Bacterial Viruses (<a href="www.phage.ulaval.ca">www.phage.ulaval.ca</a>) and stored in -80°C. In the experiments, the bacteria were cultured in TSB or BHI at 37°C. Phage stocks were prepared by harvesting the soft-agar of a confluent double-agar plate, to which four ml of media were added, followed by centrifugation (11000g) and filtration. Phage titration was made by plaque forming unit counts using the double-agar method.

# **Bioinformatic analysis**

Phage M102 <sup>22</sup> genome is publicly available (RefGenome: NC\_012884). We used HHpred <sup>23,24</sup> to analyse M102 ORFS and to find hits associated to Ig-like domains and carbohydrate binding.

### Phage binding to mucin

To evaluate phage M102 binding to mucin, BHI (Brain Heart Infusion Broth, Sigma) agar plates were prepared with or without 1% mucin in its composition. Purified porcine mucin (Sigma, catalog no. M1778) was used as source. The phage was diluted to a concentration of  $2.5 \times 10^2$  pfu (plaque forming units) ml<sup>-1</sup> in liquid BHI media and five milliliters of the dilution were added to the plates containing mucin or not. The plates were kept under agitation for 30 minutes, then the liquid was removed by careful pipetting and three milliliters of BHI soft agar containing the *S. mutans* host was added to each plate. Plates were incubated overnight at 37°C and plaques enumerated. The experiment was done in triplicates.

# Influence of mucin on S. mutans growth and biofilm formation

Influence of mucin supplementation on *S. mutans* was studied first in 1x (i.e. undiluted) TSB (Tryptic Soy Broth, Sigma) supplemented with 0.5% yeast extract and 0.5% K<sub>2</sub>HPO<sub>4</sub>). Fresh bacterial culture was inoculated into 5 ml of TSB supplemented with 0, 0.05 or 0.1 % mucin in triplicates, and cultured overnight in 37°C under constant shaking (200 rpm). OD (at 595 nm) of the cultures was measured in 6 technical replicates per tube. Data were analysed using Kruskal-Wallis non-parametric ANOVA in SPSS.

In a second experiment we explored the ability of *S. mutans* to grow with mucin without any additional energy source. Bacteria were inoculated in 5 ml of sterile H<sub>2</sub>O without mucin and with 0.2% and 0.5% mucin supplementation in triplicates. These cultures were grown in 37°C under constant shaking (200 rpm). Bacterial growth was recorded by estimating cfu (colony forming units) ml<sup>-1</sup> via applying 2 µl drops of the ten-fold dilutions of culture on BHI plates. Data were analysed with Kruskal-Wallis non-parametric ANOVA in SPSS.

Finally, we explored how different nutrient conditions and mucin concentrations influence S. mutans biofilm formation. In this experiment, overnight-grown S. mutans was inoculated 1:10 000 into sterile H<sub>2</sub>O, 0.1xBHI (i.e. ten times diluted BHI) and 1x BHI, supplemented with 0%, 0.2% or 0.5% mucin. Each treatment was done in six replicates in 150 µl volume, on a sterile 96-well plate, and incubated for 2 days at 37°C without shaking. To determine the biofilm formation, all medium was removed from the wells. Plates were washed three times with MilliQ water and stained 45 min with 0.1% crystal violet. After staining, the plates were washed again three times with MilliQ water, dried, and de-stained with ethanol for 15 minutes. OD (at 595 nm) of 100 µl samples were measured to analyse biofilm formation (Multiskan FC, Thermo Scientific, China). Statistical differences between mucin concentrations and the control were analysed using T-test in GraphPad Prism 9. 

# <u>Influence of mucin on phage-bacterium interaction</u>

First, we explored the effect of mucin supplementation and culture nutrient concentration on *S. mutans*-phage interaction in TSB medium. Overnight cultures were diluted to 1:10 000 and inoculated to 1x and 0.1xTSB supplemented with 0.5% mucin in triplicates (control cultures were done without mucin), and incubated in 37°C for 4h, after which phage M102 was added ( $10^5$  PFU ml<sup>-1</sup>). Samplings of 100  $\mu$ l were made 20h after the infection, mixed with 10  $\mu$ l of chloroform, and titrated. Another sampling was made 48h after the start of the experiment and also titrated. Kruskal-Wallis non-parametric ANOVA was used to analyse the results.

To observe the effect of mucin and phage on bacterial growth and biofilm formation, 2% porcine mucin was added to BHI to reach final concentration of 0.2%, and inoculated with 10<sup>4</sup> CFU ml<sup>-1</sup> of overnight-grown *S. mutans*. Control cultures were done in BHI without mucin. Half of the cultures were also (simultaneously) inoculated with bacteriophage M102 at MOI of 1. From each treatment, 150 µl culture was distributed into a sterile 96-well plate in six replicates, and incubated for 2 days at 37°C without shaking. After incubation, bacterial and phage numbers were sampled from 3 wells per treatment by standard plating methods. Then, the remaining biofilm from all six replicate wells was determined by crystal violet staining as explained above. The data were analysed with non-parametric ANOVA.

To understand the effect of mucosal environment on phage resistance, we co-cultured *S. mutans* with M102 phage in 0% and 0.2% mucin supplementation (in BHI, 37°C, 200 rpm) in triplicate 5 ml cultures. Samples were plated after 72 hours and bacterial colony forming units were recorded. Phage titers at the end of the experiment were determined with the original *S. mutans* host. Up to 10 bacterial colonies were picked from each replicate and purified by three rounds of colony-picking and plating. Purified colonies were then inoculated into fresh BHI, grown at 37°C with shaking (200 rpm), and preserved into -80°C freezer stocks with 10% glycerol. Phage resistance of the isolates was determined with a plaque assay using the ancestral M102 phage.

### **RESULTS**

# Phage M102 adhesion to mucin-containing plates

A brief survey of the M102 genome (NC\_012884) revealed that it has a putative carbohydrate-binding domain in ORF13 (HHPred <sup>25</sup>: Lactobacillus phage J-1 carbohydrate binding module, Probability: 99.95%, E-value: 8.8e-26) as a tail component, which may mediate binding to mucin, or to the bacterial host. We evaluated the ability of M102 to bind *in vitro* to purified mucin. Such approach has been applied before to evaluate mucin-binding capacity of different

phages  $^{6,10}$ . Despite a trend in having slightly more phages in mucin containing plates (on average 1.48 more phages on mucin plates), there is no significant difference between the tested conditions (Figure 1, unpaired t test, p= 0.1310).

# Influence of mucin on S. mutans growth and biofilm formation

We measured the effect of mucin supplementation on *S. mutans* growth in 5 ml triplicate cultures in TSB alone or supplemented with mucin. Mucin supplementation significantly influenced bacterial growth (Kruskal-Wallis non-parametric ANOVA, test statistic=47.167, df =2, p<0.01), measured as OD of the cultures. In pairwise comparisons all treatments significantly differed from each other (p-values Bonferroni-corrected) (Figure 2A).

When the bacteria were inoculated in pure water, there was no bacterial growth observed without mucin supplementation, whereas 0.2% and 0.5% mucin resulted in numbers of  $7.2\times10^5$  and  $1.1\times10^5$  CFU ml<sup>-1</sup>, respectively (Figure 2B). There was no statistical difference between these two treatments.

Although it was evident that mucin supplementation benefits S. mutans growth, this was not reflected in biofilm formation (Figure 3). Mucin presence in water cultures did not lead to biofilm formation, and the same was seen when diluted (0.1x) medium was used. In complete (1x) medium, biofilm was detected in all conditions, with or without mucin. However, t-tests revealed that mucin addition significantly decreased biofilm formation in 0.1% BHI (p<0.001 0% vs 0.2% mucin and 0% vs 0.5%) and in the 1x BHI (p<0.001 for 0% vs 0.2% mucin and 0% vs 0.5%). It may be that the presence of nutrients (culture media) favours biofilm formation while mucin could serve as an inducer of the planktonic state.

# Influence of mucin on phage-bacterium interaction

First, we explored the effect of mucin supplementation on *S. mutans*-phage interaction in TSB medium. Twenty hours after phage inoculation there was a significant increase in phage titer when 0.1 TSB was supplemented with 0.5% mucin (Kruskall Wallis 1-way ANOVA p=0.0094, bonferroni-corrected p-value for pairwise comparisons 0.027) (Figure 4). After 48h, phage titers were similar in all treatments (data not shown).

To understand whether the improved phage growth was because of mucin presence or increased bacterial population in the mucin condition, we cultured *S. mutans* in presence of phage M102 and mucin for 48h in 1x BHI. In the non-infected cultures bacteria titers did not differ between control and mucin conditions. However, bacterial numbers significantly differed between treatments (Kurskal-Wallis 1-way ANOVA; test statistic=9.359, df=3, p=0.025) in the presence of the phage (Figure 5). Notably, phage significantly reduced bacterial numbers only when mucin was present. Phage titration of the treatment indicated, that phage replicated only in presence of 0.2% mucin (titer 10<sup>7</sup> pfu ml<sup>-1</sup> vs 10<sup>3</sup> pfu ml<sup>-1</sup> in treatment without mucin addition). Phage presence also significantly affected biofilm formation, but only when mucin was present (Figure 5B).

#### Effect of mucin on phage-bacterium interaction and phage resistance

To understand the effect of mucosal environment on phage resistance, we co-cultured *S. mutans* with phage M102 in 0% and 0.2% mucin supplementation (in BHI, 37°C, 200 rpm) in triplicate 5 ml cultures. Compared to the previous experiment, we used a longer incubation time to allow phage resistance to evolve. After 72 hours, bacterial colony forming units were recorded and phage titers determined with the original *S. mutans* host (Figure 6). Ten bacterial colonies per

replicate were pure cultured from treatments evolving with the phage (and 3 colonies per replicate from phage-free conditions) and their phage resistance determined by plaque assays. Under mucin supplementation and phage exposure, 26 of 30 colonies showed a reduced susceptibility to M102 after 72 h. All colonies isolated from the control conditions remained sentisitive to phage infection.

Bacterial densities did not statistically differ between experimental conditions but, again, phage replication was detected only in 0.2% mucin (average  $1.05 \times 10^7$  PFU mL<sup>-1</sup>). In replicates without mucin no plaques were detected. Since also the phage-only control was negative for phage plaques, the phage may not have survived the 3-day culture conditions, or the titer was under the detection limit.

#### **Discussion**

S.mutans causes tooth decay i.e. dental caries, a common condition which is hard to treat. The challenges of the increasing antibiotic resistance in general and the susceptibility of the aging population to infections require developing approaches that target bacterial pathogens also in the oral context. This has led to a growing interest towards phage therapy. Phages against S. mutans have already been shown to be potential preventive agents against caries in vitro and in vivo models, highlighting its relevance as a non-invasive treatment <sup>26</sup>. Therefore, we need to understand how phages infect bacteria and how bacteria resist phage infections in the mucosal environment.

To our knowledge, there has not been prior studies addressing *S. mutans* phage infections in presence of mucins. In general, phage-bacterium interactions are most often explored in standard defined growth media, where the possible signals from the vertebrate host targeted by the bacteria are absent. However, in real life, the interactions between pathogenic bacteria and phages often occur on the mucosal surfaces of vertebrate hosts. Our results show that mucosal environment (achieved by mucin supplementation of growth media) significantly alters *S. mutans* growth, phage sensitivity and phage resistance. These results underline the importance of understanding the effect of mucosa on phage-bacterium interactions.

Our results indicate that *S. mutans* can benefit from mucin as an additional energy source, as mucin supplementation increases bacterial replication at least if added in TSB (Figure 2), where additional nutrients are available for growth. This is in line with results obtained previously<sup>7</sup>, where addition of mucin and glucose enhanced *S. mutans* growth in chemically defined medium. Furthermore, presence of mucins increases *S. mutans* survival <sup>17.</sup> This may explain the higher bacterial numbers in mucin-supplemented H<sub>2</sub>O observed in our study in contrast to none in replicates without mucin (Figure 2b). Yet, it is clear that the interactions between *S. mutans* and mucins are more complex *in vivo*. For example, the microbial diversity found in the oral environment can influence *S. mutans* behaviour and phenotype. Mucins have been shown to influence microbial interactions and support the coexistence of *S. mutans* and its competitors <sup>27</sup>.

Interestingly, mucin decreased *S. mutans* biofilm formation, which was more dependent on the concentration of nutrients. A previous study using human salivary mucins suggested that mucin decreases *S. mutans* surface attachment by inducing planktonic growth form <sup>27</sup>. This may be linked directly to sensitivity to phage, as bacteria growing in the planktonic phase are not protected by biofilm, allowing phage infections. Furthermore, presence of oxygen has been shown to decrease biofilm formation in *S. mutans* <sup>19</sup>, which may be one of the reasons why the

biofilm formation was low in general in our experiments, which were made in aerobic conditions. The influence of mucin under anaerobic conditions is a factor to be considered in future research.

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Our results show that mucins significantly influence S. mutans phage survival and replication, exemplifying the role of mucosal surfaces on the phage-bacterium interaction and potentially phage therapy approaches. Studies comparing the density of free and metazoan-associated phages have found that phages tend to concentrate on mucosal surfaces <sup>10</sup>. For example, 10<sup>8</sup> virus-like particles per ml can be found in the human gut microbiota, but the density is higher in the gut mucosa (10<sup>9</sup> phages per ml)<sup>28</sup>. This can be explained by the affinity of some phages towards mucins: some tailed dsDNA phages have (structural) protein domains that mediate their adherence to mucin glycoproteins found in the mucosal surfaces. A bioinformatic analysis of 246 dsDNA tailed-phage genomes suggested that roughly 25% have proteins with Ig-like folds, all related to the viral structure <sup>29</sup>. More recently, also other domains <sup>30, 31</sup> have been found and suggested to facilitate phage binding to mucosa. Phage M102 22 has a putative carbohydrate-binding domain in ORF13 which might mediate the phage-mucus interaction, but the role of this domain remains unknown so far. Nevertheless, although M102 mucin binding in vitro was not significant, the ratio between plaque count in mucin divided by plaque count in control was 1.48. Since it has been shown that phage-mucin interactions are transient but constant <sup>20</sup>, even a slightly positive ratio allied to the numerous interactions between the phage and mucins may still play a role in maintaining phages in the mucosa. Furthermore, while the phage titers in mucin-free cultures were below detection limit, it is also possible that presence of mucins stabilize phage particles, influencing their survival in the mucosal environment.

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While phage-host interactions in S. mutans have been previously explored <sup>26,32,33</sup>, there is a lack of knowledge on how the mucosal environment influences bacterial susceptibility to phage. We observed that mucin has a major role for phage susceptibility: mucin supplementation caused a 4-log increase in phage titer of S. mutans. While the exact mechanisms for this phenomenon are unknown, it is possible that phages benefit from higher bacterial replication or that they exploit their changed metabolic state. It is also possible that the upregulation of bacterial virulence factors (related to cell wall structures, secretion, metabolism or motility) in the mucosal environment increases the availability of phage receptors. This could explain the previously observed increased virulence of pathogens in the mucosal environment 4-6. In our experiments, it seems evident that the choice of medium influences the impact of mucin on bacterial growth and phage sensitivity. In TSB, which is a less rich medium than BHI, phage replication was improved only in the diluted medium (Figure 4) whereas in BHI phage replication (and a drop in bacterial population density) was observed only under mucin supplementation. Yet, in BHI, mucin supplementation did not cause any differences in bacterial numbers in absence of phage (Figure 5). This suggests that mucininduced expressional changes are more likely to cause sensitivity to phage infection. Consequently, phage resistance was elicited only under mucin supplementation when the bacteria were co-culture with phage for 72h. This is seen as a comparable bacterial population size in phage-treated and control culture in Figure 6a. While phage resistance in S. mutans has been observed in previous studies without mucin <sup>32,33</sup>, it remains to be studied how the mucosal environment influences the evolution of resistance in S. mutans, e.g. CRISPR-Cas <sup>18</sup>.

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The influence of mucins on bacterial phage sensitivity has also been studied previously e.g. with *Salmonella enterica* <sup>34</sup>, and many other bacterial species <sup>35</sup>. However, in some cases mucins may also impair phage infectivity <sup>6,17</sup>. Such phages are probably more efficient in

targeting pathogens outside the mucosa. Differences in the capacity to adhere to mucus (or to replicate in mucosal environment) could be considered as one additional phenotypic feature of phage-bacterium interaction, that could be exploited in building phage cocktails for therapy purposes. A combination of phages with different strategies (mucin-binding or not) may produce efficient treatment and prevention tools that target pathogens both from the infection site and during transmission.

The mucin concentrations used in these experiments are relevant and comparable to what is observed in the body, although mucin species and concentrations vary depending on body site and and in response to physical conditions, disease and age <sup>36–39</sup>. For example, in the gastric environment the concentration of the mucins is ~5% <sup>21</sup>, but the density of the mucosal layer is not consistent, as the mucin concentration is highest near the cell surface and less dense at the outer layer. The oral mucosal environment contains several different mucins; secreted salivary mucins MUC5B, MUC7 and MUC19, and membrane-associated mucins MUC1 and MUC4 <sup>40</sup>. The salivary mucin levels are generally higher in patients with dental karies, e.g. with MUC5B concentration varying from 0.06 to 2.34 ng mL<sup>-1</sup> <sup>38</sup>. While the exact mucin species composition of the porcine mucin used in our experiments is not known, it mainly consists of MUC2. It thus seems that *S. mutans* may not have a specific affinity towards certain mucins as observed in some other bacteria <sup>40</sup>. Furthermore, it is important to note that porcine mucin has been shown previously to be relevant for phage-bacterium studies in also other species, eliciting a similar increase in phage replication<sup>6</sup>.

 Combined with previous research, these results highlight the effect of mucosal environment on phage-bacterium interactions, and emphasizes the ecological relevance of microbial interactions in the eukaryotic mucosa, with implications for the outcome and prevention of diseases.

#### **Conflict of Interest**

G.M.D.F.A., L.-R.S. and University of Jyväskylä have patented the commercial use of mucin in a patent titled "Improved methods and culture media for production, quantification and isolation of bacteriophages" (FI20185086, PCT/FI2019/050073).

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# **FIGURES**

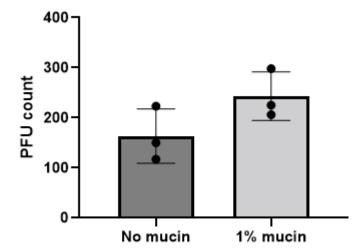


Figure 1. Streptococcus mutans phage M102 adhesion to mucin. Each data point represents an individual replicate that consists of phage plaques count in an agar plate. The mean and standard deviation of each condition is indicated in the graph.

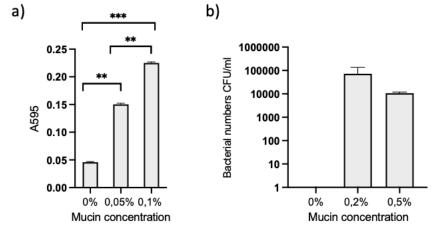


Figure 2. Effect of mucin supplementation on Streptococcus mutans growth. a) Mean optical density (+/- Standard error) of bacteria grown in TSB supplemented with 0%, 0.05% and 0.1% ucin, and b) mean colony forming units per ml (+/- S.E.) in bacteria grown in sterile water supplemented with 0%, 0.2% and 0.5% mucin. Asterisks indicate statistical significance (p-values: \*\*\*< 0.001, \*\*=0.001-0.01.\*=0.01-0.05)

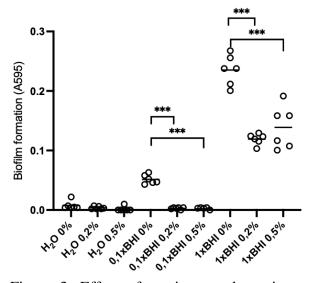


Figure 3. Effect of nutrients and mucin concentration on *Streptococcus mutans* biofilm formation. Bacteria were cultured without nutrients  $(H_2O)$ , in 0.1X (ten times diluted) and 1X (normal) BHI under 0%, 0.2% and 0.5% mucin supplementation for 48h. Asterisk indicates statistical significance (\*\*\*= p-value <0.001)

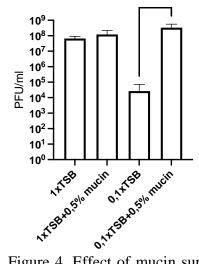


Figure 4. Effect of mucin supplementation on M102 phage titers in TSB (1X: normal TSB, 0.1X: 10 times diluted TSB). Asterisk indicates statistical significance (\* = p-value <0.05)

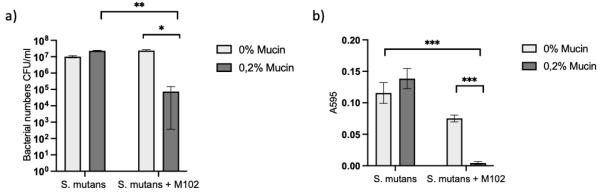


Figure 5. S. mutans growth and biofilm formation when cultured with 0.2% mucin supplementation and phage M102 for 48 hours. A) Mean bacterial numbers +/- SE, b) bacterial biofilm formation (mean OD+/- SE). Asterisks indicate statistical significance between treatments (p-values: \*\*\*< 0.001, \*\*=0.001-0.01.\*=0.01-0.05)

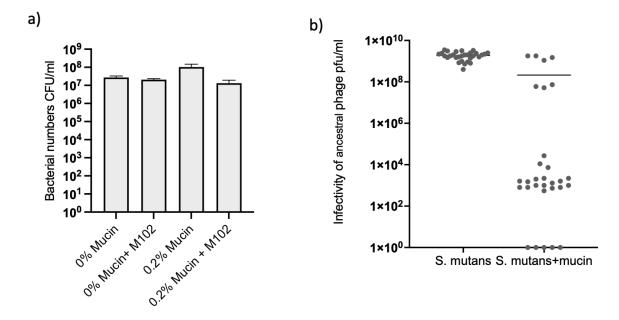


Figure 6. Phage resistance experiment in mucosal environment. Phage and bacteria were cocultured for 72h to allow phage resistance to evolve. A) Bacterial cell numbers with phage M102 and 0,2% mucin supplementation. B) phage infectivity (pfu/ml by plaque assay) of isolates originating from phage exposure with and without mucin supplementation.