

# Deciphering the role of monosaccharides during phage infection of Staphylococcus aureus

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- 3 Graphical table of contents. S. aureus cell wall was deciphered using a phage-decorated atomic force
- 4 spectroscopy tip in order to identify major receptors involved in phage adhesion and infection of host cell.

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7	phage infection of <i>Staphylococcus aureus</i> .
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# 16 ABSTRACT

17 As phages are extensively investigated as novel therapy tools but also as transfer agents for antibiotic 18 resistance genes, thorough understanding of phage-host interactions becomes crucial. Prerequisite for 19 phage infection is its adhesion to the host surface. Here, we used atomic force microscopy-based single-20 particle force spectroscopy with phage-decorated tips to decipher the adhesion of phage 187 on living 21 Staphylococcus aureus cells. We found that addition of free N-acetyl-D-glucosamine was able to decrease 22 phage adhesion, suggesting that this monosaccharide plays major role in phage 187 infection of S. aureus. 23 Moreover, phage 187 adhesion on monosaccharide-coated model surfaces combined with plaque forming 24 unit counts suggested that a direct link can be established between the propensity to bind to a saccharide 25 and the capability of the latter to inhibit phage infection. On a nanoscale level, single-particle force 26 spectroscopy was successfully used to identify a major receptor required for phage 187 infection of S. 27 aureus but also evidenced that this receptor was responsible for phage adhesion on host-cells. Our work 28 demonstrates that single-particle force spectroscopy is a powerful platform to screen and predict the 29 molecular target of phages on their host surfaces.

30

#### 31 KEYWORDS

Atomic force microscopy, Single-particle force spectroscopy, Phage 187, S. aureus, Phage-bacteria
 interactions

# 35 1. INTRODUCTION

36 Phages are viruses that infect specific bacteria and that can undergo two different life cycles: lysogenic 37 and lytic [1-3]. Lysogenic phages are known to be one of the causes for the increase of multi drug resistance 38 as they promote horizontal gene transfer (HGT) among bacteria [4]. Lytic phages, on the other hand, are 39 suggested to be one of the solutions for the raise of antibiotic-resistant pathogens with phage therapy as 40 they can infect and rapidly kill these pathogens [5-8]. Phages are commercially available as biocontrol 41 agents to help prevent colonization by pathogenic bacteria in agriculture, livestock and food production 42 [9-11], and they present potential applications in wastewater treatment as a method of controlling 43 pathogens in sewage water [12, 13]. After being eclipsed by the use of antibiotics for decades, phages have 44 been regaining attention in the last years due to their implications in the raise of antibiotic-resistant 45 infections and because of their vast potential of applications as antimicrobial agents.

46 All viruses including phages start their infection cycle by non-specific and specific interactions at the 47 surface of their host cells [3, 14]. The recognition of the specific receptor is a key point in the host spectrum of a phage. While very little is known about non-specific interactions, some specific phage receptors are 48 49 known, but again knowledge remains limited. Recognition is achieved by phage receptor binding proteins 50 (RBPs) and is dependent on location of their target receptors on the bacteria wall surface as well as 51 chemical and structural peculiarities [3, 14-17]. RBPs have strong affinities for specific receptors and allow 52 phages to anchor on their host. Phages have single- or double-stranded DNA or RNA genome surrounded 53 by a capsid. For the vast majority of phages (around 96%), the capsid (or the head) is connected to a protein 54 tail that has RBPs at its other end [18]. Once anchored onto the cell surface, phages can inject their genetic 55 material inside the host cell by using peptidoglycan hydrolases to degrade cell wall locally without lysing 56 the bacteria. Host cells provide the molecular building blocks and enzymes required to replicate phages 57 and produce progeny [1-3].

58 Understanding and characterizing phage-bacteria interactions is critical as they dictate infection [15]. In 59 parallel of genomic and microbiology characterization, physico-chemical characterization of phages gave 60 decisive insights notably through atomic force microscopy (AFM) which contributed intensively in imaging 61 and characterizing the infection process of phages [19-21]. AFM-based force spectroscopy was revealed 62 to be a powerful tool to characterize bacteria interactions with the use of single-molecule force 63 spectroscopy (SMFS) and single-cell force spectroscopy (SCFS) [22-26]. For instance, AFM has already been 64 used to decipher functions of viral attachment proteins in the initiation of infection with reovirus-65 decorated tips by SMFS on living cells [27]. AFM enables in situ characterization of living bacterial cells 66 interactions with phages. However, most studies on living cells focus on the effects of phage infections on 67 bacteria but very few studies focused on phage interactions themselves with single living host-cells to 68 investigate adhesion [21, 28, 29]. In this work, we focus on specific recognition of living Staphylococcus 69 aureus cells by phage 187 which belongs to the Siphoviridae family [30]. S. aureus, a Gram-positive 70 bacterium, is a human pathogen of major concerns that can colonize a large variety of tissues in the body 71 [31]. It commonly causes superficial and invasive infections leading to severe morbidity and mortality [32]. 72 Treatments of *S. aureus* infections rely on long-term antibiotics which led to the raise of resistant strains 73 such as methicillin-resistant S. aureus (MRSA) [33]. In Gram-positive bacteria, lipoteichoic acid (LTA) along 74 with wall teichoic acid (WTA) are two of the major components of the cell envelop and are critical for 75 growth, survival and pathogenicity [34, 35]. LTA and WTA form a polyanionic barrier that is required to 76 maintain metal cation homeostasis and assist in transfers of ions, nutrients and proteins [36-38]. 77 Peptidoglicans (PG) in the cell wall form a network on which polymers like WTA, polysaccharides and 78 proteins are covalently linked. S. aureus WTA is generally made of repeating units composed of 1,5-D-79 ribitol-phosphate (RboP) tailored with D-alanine esters and mono- or oligosaccharides, commonly N-80 acetyl-D-glucosamine (GlcNAc) or N-acetyl-D-galactosamine (GalNAc) in certain cases [35, 39-41].

81 In the present study, AFM-based single particle force spectroscopy (SPFS) was used to map and probe 82 interactions between single living S. aureus cells and phage 187 (Figure 1a). Effects of addition of free 83 monosaccharides on phage infectivity were investigated as well as phage interactions with different 84 monosaccharide-coated model surfaces. Because of S. aureus wall composition, GlcNAc and GalNAc were 85 selected as the investigated monosaccharides along with their non-amino homologues: glucose (Glc) and 86 galactose (Gal). This study provided valuable insights into the correlation between the specific ability of 87 phage 187 to adhere on monosaccharide-coated surfaces and the capability of the monosaccharides to 88 inhibit phage infection. On a nanoscale level, SPFS was used to identify a major receptor responsible for 89 phage adhesion on host-cells and thus, a major receptor for phage infection initiation. Our findings helped 90 building a predictive model designed to select best suitable saccharides to reduce phage infectivity by 91 screening saccharide-coated model surfaces with phage-decorated AFM probes. Models developed with 92 SPFS can also be used to predict molecular targets of phages on their host surfaces. This study expands 93 the repertoire of AFM-based force spectroscopy and provides insights into correlation between phage 94 adhesion and infectivity.

# 95 2. MATERIALS AND METHODS

#### 96 **2.1. Bacteria strains and phage 187.**

97 Staphylococcus aureus phage 187 (HER 239) and Staphylococcus aureus strain 187 (HER 1239) were 98 provided by Félix d'Hérelle Reference Center for Bacterial Viruses from the Université Laval (Québec, 99 Canada). Phages were purified as previously described [42]. Briefly, after replication, phage 187 100 suspensions were centrifugated at 4,000 g for 20 min at 4°C and the supernatant was filtered using a 0.45 101 µm membrane (Millex<sup>®</sup>-GP, Millipore). Then 30 mL of this suspension was concentrated in 10 mL by 102 Amicon Ultra-15 (PLHK, membrane Ultracel-PL, 100 kDa) by centrifugation at 1,000 g for 3 min. The phage 103 suspension (2.5 mL) was purified by cesium chloride gradient after addition of 3.88 g CsCl (final volume: 6 104 mL) and centrifugation at 113,000 q for 18h at 15°C. About 1.5 mL of phage suspension was extracted from 105 central strip and then purified by dialysis (100 kDa molecular weight cutoff, Spectrum) against phosphate 106 buffered saline (0.1X) for 14 h. The concentration of phages 187 obtained following the above protocol 107 reached 10<sup>9</sup> PFU/mL. S. aureus was routinely grown at 37°C in tryptic soy broth (TSB) media. For phage 108 infection, 10<sup>2</sup> PFU/mL of phage 187 was incubated with or without free monosaccharides (final 109 concentration 10 mM) in Tris Buffered Saline supplemented with 1 mM CaCl<sub>2</sub> and MgSO<sub>4</sub> during 2h at room 110 temperature under gentle agitation. S. aureus was cultivated in complemented tryptic soy broth to log-111 phase (OD<sub>600</sub>, approximately 0.2) and infected with equal volume of phage 187. Phages were allowed to 112 absorb for 5 min on *S. aureus* before plating on complemented tryptic soy agar (TSA) for titration.

113

#### 2.2. Surface and AFM tip functionalization with phages.

114 Phages were attached covalently on self-assembled monolayers (SAM) of carboxyl-terminated 115 alkanethiols. Briefly, gold substrates and gold coated NPG-D (Bruker) AFM tips were immersed overnight 116 in a solution of 0.1 mM 16-mercaptododecahexanoic acid and 0.9 mM 11-mercapto-1-undecanol to form 117 the SAM. Then, they were washed with ethanol, dried under N<sub>2</sub> flow and carboxylic groups were activated 118 by immersion 30 min in a solution of N-hydroxysuccinimide (NHS, 20 mg/mL) and 1-ethyl-3-(3-119 dimethylaminopropyl)-carbodiimide (EDC, 50 mg/mL). After rinsing with ultrapure water, 2 x 10<sup>8</sup> PFU of 120 purified phages were added for 1 hour. NHS/EDC coupling formed strong covalent amide bonds between 121 the carboxylic group of the 16-mercaptododecahexanoic acid thiol present on the functionalized goldcoated tips and the amine groups from the phage surface. Finally, surfaces and tips were rinsed in TBS 122 123 buffer containing 1 mM CaCl<sub>2</sub> and MgSO<sub>4</sub> (TBS<sup>+</sup>) and stored for max. 48 h in the same buffer until their use. 124 For AFM images in air, surfaces were gently rinsed in 3 baths of ultrapure water and let to dry overnight 125 at 30°C before imaging.

126

# 2.3. AFM imaging and force spectroscopy.

All experiments were achieved at room temperature using a Bioscope Resolve AFM (Bruker corporation,
Santa Barbara, CA) and cantilevers spring constants were determined by the thermal noise method [43].

129 Phage immobilized on substrates were imaged in peak-force tapping mode using HIRS-F-B tips in air or in 130 TBS<sup>+</sup>. For force measurements on cells, stationary phase cultures of *S. aureus* were centrifuged, washed 131 twice in TBS<sup>+</sup> and filtered into porous polycarbonate membranes (Millipore). After filtration and rinsing, a 132 1 x 1 cm<sup>2</sup> piece of membrane was cut and attached with double sided adhesive tape at the bottom of a 133 Petri dish and immersed in TBS<sup>+</sup>. First, bare SNL-D tips were used to localize and image individual cells. 134 Then, the tip was replaced by phage decorated tip and adhesion maps of 32 x 32 FD curves on areas of 500 135 x 500 nm<sup>2</sup> were recorded on the cell surface with 250 pN applied force, 1  $\mu$ m/s constant approach and 136 retraction speeds and 250 ms contact time. For inhibition tests, 10 mM of free monosaccharides were 137 added to the Petri dish and adhesion maps were subsequently recorded on the same cells. For force 138 spectroscopy on monosaccharide-coated model surfaces, silicon substrates were immersed overnight in a 139 mixture of 0.9 mM tetraethoxysilane (TEOS, Si(OCH<sub>2</sub>CH<sub>3</sub>)<sub>4</sub>), and 0.1 mM isocyanatopropyltriethoxysilane 140 (ICPTES, (C<sub>2</sub>H<sub>5</sub>O)<sub>3</sub>Si(CH<sub>2</sub>)<sub>3</sub>NCO) for silanization and functionalization with isocyanate moieties. Then, they 141 were washed with ethanol, dried under N<sub>2</sub> flow and immersed in 10 mM GlcNAc, GalNAc, Glc and Gal for 142 24 hours. Monosaccharide coated substrates were then rinsed with TBS<sup>+</sup>, imaged in contact mode with 143 bare SNL-D tips and probed with phage decorated NPG-D tips by recording adhesion maps of 5 x 5  $\mu$ m<sup>2</sup>.

144 3. RESULTS AND DISCUSSION

145 3.1. Phage 187 infects S. aureus and can be grafted on gold-coated AFM cantilevers.

The first step of this study was to evaluate the infectivity of phage 187 on *S. aureus* by phage plaque assay. Briefly, lysis plaques are clear zones that appear in a homogenous bacteria culture on agar medium due to lysis by phages. Presence of lysis plaques shows the ability of the phage to infect and lyse the bacteria. Lysis plaques are the result of cycles of infection due to one phage (or one aggregate of phages) and occurring cell by cell in all directions from the original infection center thus producing circular areas of lysis called plaque forming unit (PFU) [44]. They represent a simple, cost-efficient and fast technique to

evidence and quantify phage infection [45]. When *S. aureus* cells were in contact with purified phages 187,
lysis plaques of ~1 mm in diameter were observed (Figure 1b).

154 To probe the interactions between phage 187 and S. aureus, we hypothesized that phages can be 155 covalently grafted on gold coated AFM tips using N-hydroxysuccinimide/N-(3-dimethyl-aminopropyl)-N'-156 ethylcarbodiimide hydrochloride (NHS/EDC) coupling. To confirm this hypothesis, we first functionalized 157 and imaged flat gold substrate. Figure 2 shows topography images of phages covalently attached on gold-158 coated substrates. Phage 187 is known to have polyhedral shaped heads connected to a tubbed-like tail 159 [46]. In air, the typical shape of Siphoviridae was confirmed with the presence of polyhedral heads linked 160 to tails (Figure 2a). Imaging single phage at high resolution revealed a full length of ~200 nm with a head 161 of ~65 nm from the apex to the head-tail joint and a tail length of ~140 nm (Figure 2b). Head height was 162 estimated at ~10 nm and the tail revealed a tubbed-like shape of around 6 nm in height. These 163 morphological features are in good accordance with previous observations of phage 187 achieved by 164 electron microscopy [46]. At the distal end of the tail, an increase of height was observed. We associated 165 it to the baseplate of the phage to which RBPs are attached as already suggested with other phages 166 observed by AFM in the literature [47]. In liquid, the head height was ~25 nm (data not shown). We 167 attributed this height difference to particle collapsing during air drying which is a common phenomenon 168 in AFM phage observations [47]. Remarkably, whereas phage heads were successfully imaged in liquid, 169 tails were not detected or appeared as poorly defined streaks (Figure 2c) probably because they were too 170 flexible or too mobile and were pushed by the scanning tip. As phages were successfully observed in liquid 171 and air, we concluded that phage 187 can be successfully attached to gold substrates. Phages grafted on 172 gold-coated substrate using NHS/EDC coupling led to strong covalent amide bonds that resist forces 173 superior to 2 nN [48, 49]. Therefore, as long as the forces measured by SMFS do not exceed the 174 nanonewton, the amide bond between the phages and the gold-coated AFM tips was considered covalent 175 and irreversible. Thus, we decided to use this approach to functionalize gold-coated AFM tips with phages

in order to measure their interaction with living *S. aureus* cell surfaces by mean of AFM-based SPFS (Figure2d).

178

# 3.2. Phage 187 recognizes *S. aureus* cell wall with high adhesion frequency.

179 We probed single living S. aureus cells with phage-decorated AFM tips. Figure 3 and S1 show adhesion 180 maps, adhesion force and rupture length histograms as well as representative force-distance (FD) curves 181 obtained by SPFS on live bacteria. For all the cells, adhesion maps did not show cluster patterns, *i.e.* 182 adhesive pixels were randomly spread on the maps (Figure 3a and S1a). This suggested that phage 183 receptors were homogeneously distributed on the host cell surface. Most of the FD curves (51% to 91% 184 depending on the cell) showed adhesive peaks except for cell#S3 (Figure S1b-c) that showed an adhesion 185 frequency of only 17%. Variations in adhesion frequencies between cells were suggested to be caused by 186 slight heterogeneity in cell wall microarchitecture, and composition and/or the angle and position of 187 phage-decorated AFM cantilevers during measurements. Forces mainly ranged from 50 pN to 600 pN 188 (Figure 3b and S1b). This confirmed that phage-bacteria interactions forces were inferior than typical 189 covalent bond forces that exist between the phages and the AFM tips [48]. Thus, we considered that during 190 the repeating approaches and retractions of the AFM tips throughout measurements, weak bonds 191 between phages and cells ruptured before covalent bonds between the phages and the tips. This was 192 further confirmed by the consistent adhesion signatures obtained throughout all the measurements in 193 Figure 3 and S1 and also confirmed with the homogeneousness of the adhesion maps obtained. The 194 majority of rupture lengths ranged between 10 nm and 300 nm (Figure 3c and S1c). Representative FD 195 curves are presented in Figure 3c and S1c as examples. They show the retraction of the AFM tip decorated 196 with the phages. Some curves did not present any adhesion peaks and were considered as "non-adhesive 197 events". By contrast, adhesive FD curves presented a single or multiple adhesion peaks that we attributed 198 to the interactions between the phage RBPs and the receptors on the cell wall.

## 3.3. SPFS reveals major cell wall receptors for phage adhesion.

200 Phage 187 showed homogenous recognition of S. aureus cell wall suggesting phage receptors are 201 distributed evenly on the surface of cells. S. aureus WTA, which is one of the most abundant components 202 in cell wall, generally presents repeating units composed of RboP with mono- or oligosaccharides residues 203 like GlcNAc or GalNAc [35, 39-41]. We used SPFS to decipher phage interactions with GlcNAc and GalNAc 204 in order to conclude if these monosaccharides lead to specific adhesion of phage 187. Specific adhesion 205 would suggest a probable implication of these molecules in phage recognition process. Interactions with 206 non-amino equivalents, Glc and Gal, were also deciphered. Effect of different monosaccharides on phage 207 infectivity was first investigated at the population scale by incubating phage 187 with free 208 monosaccharides and then quantifying infectivity by PFU counting (Figure 4). When  $1 \times 10^2$  PFU of phage 209 187 was put in contact with free Glc, Gal, and GalNAc for 2h prior to infection, total phage-infected S. 210 aureus cells decreased respectively of 22%, 12% and 37% compared to the absence of free 211 monosaccharide. Remarkably, total inhibition was observed with free GlcNAc as no lysis plagues were 212 observed suggesting that no infection occurred or infection was below the detection limit (Figure 4). This 213 suggests that phage 187 RBPs were already saturated with free GlcNAc residues and thus were unable to 214 adsorb and anchor to S. aureus cells. Therefore, infection process could not be initiated resulting in a total 215 inhibition of infectivity.

To determine the effect of monosaccharides on phage adhesion at the single particle level, living cells were probed with AFM tips decorated with phage 187 in absence or presence of free monosaccharides. As previously discussed, slight quantitative heterogeneity in adhesion frequencies appeared that may be caused by heterogeneities in the cell population. To alleviate this possible undesired parameter and in order to compare the effect of monosaccharides on phage adhesion, the same cell was probed for each measurement before and after monosaccharide addition (Figure 5). *S. aureus* cells were immobilized in porous membranes (Figure 5a). Figure 5b shows the adhesion force histograms obtained without 223 monosaccharide and Figure 5c displays the adhesion force histograms for the same cells with different 224 monosaccharides supplemented in the media. In absence of free monosaccharides, frequencies of 225 adhesive events ranged from 69% to 82% (Figure 5b). When free monosaccharides were added to the 226 media, a decrease in adhesion was observed and frequencies of adhesive events ranged from 25% to 61% 227 (Figure 5c). For all conditions, the number of phage adhesive events on cells decreased but the intensity 228 of interactions, *i.e.* the range of adhesion forces, stayed the same (Figure 5b vs 5c). Adhesion frequency 229 decreased from 76% to 42% (i.e. a delta of 34%) when free Glc was added in media (Figure 5). With the 230 presence of free GlcNAc, a decrease of adhesion was also observed from 82% to 25% (*i.e.* a delta of 57%). 231 This shows that phage 187 adhesion on *S. aureus* was less prone to happen in presence of free GlcNAc 232 than in presence of free Glc, suggesting that the acetyl amino group of GlcNAc plays a role in phage 233 adhesion and recognition. In presence of free GalNAc, a delta of 33% of adhesive events was observed 234 (from 72% with no GalNAc to 39% with GalNAc). However, the addition of free Gal in the media showed 235 almost no effect on phage adhesion with a variation of adhesion frequency from 69% without Gal to 61% 236 with free Gal (*i.e.* a delta of 8%). Decrease in phage adhesion was more important with free GalNAc 237 compared to free Gal suggesting once again that the acetyl amino group plays a role in phage adhesion 238 and recognition. To confirm these observations, SPFS was repeated on multiple cells. Moreover, the 239 concentration of monosaccharide used is not known to cause any damages to cell walls [50]. But to 240 alleviate any possible contribution of the monosaccharides in the alteration of adhesion signatures and 241 cell integrity, control cells were imaged before and after monosaccharide addition. Addition of 242 monosaccharides did not alter the morphology of bacterial cell wall (Figure S3) thus we assumed that 243 monosaccharides did not cause any damage to the cells that could have altered the adhesion signatures. 244 Figure 6 regroups the results obtained with four cells per monosaccharide tested. GlcNAc induced once 245 more the most important adhesion decrease with a mean delta of 53% ± 4% whereas adhesion decreased 246 by only  $15\% \pm 6\%$  with free Glc. GalNAc and Gal led to a decrease of  $24\%\pm6\%$  and  $9\%\pm2\%$ , respectively.

This confirmed that phage 187 had lessened abilities to adhere on cell wall surface when free GlcNAc,
GalNAc or Glc are added in the media and that the acetyl amino groups of GlcNAc and GalNAc play a role
on phage adhesion.

# 3.4. Monosaccharide-coated model surfaces coupled with SPFS can help predict monosaccharide effects of phage inhibition.

252 In order to confirm the specific interaction of phage 187 with different monosaccharides and exclude 253 possible effects of other cell wall components, model surfaces coated with monosaccharides were 254 developed. Model surfaces coated with GlcNAc, GalNAc, Glc and Gal were probed with phage-decorated 255 tip to screen the affinity of phage 187 for each monosaccharide independently (Figure 7a). Topography 256 images of the coated surfaces revealed smooth and homogeneous deposition for all samples with a 257 thickness of ~1.3 nm (Figure 7b). Phage 187 presented high adhesion frequency towards GlcNAc model 258 surfaces (Figure 7c). Adhesion events represented around 93% of the total FD curves for the surface coated 259 with GlcNAc whereas for GalNAc, Gal and Glc, only 32%, 16% and 14% of the FD curves corresponded to 260 adhesive events, respectively. Monosaccharide containing acetyl amino groups led to higher adhesion 261 frequencies than homologous molecules lacking acetyl amino groups. Phage 187 had strongest affinities 262 with these monosaccharides (GlcNAc and GalNAc) which is consistent to the inhibition results of phage 263 187 infectivity in presence of the free monosaccharides. Figure S2 shows three independent SPFS 264 acquisitions on GlcNAc-coated model surfaces. Similarly to acquisitions on cells, adhesion maps on model 265 surfaces revealed an homogeneous distribution of adhesion events (Figure S2a). Adhesion forces ranged 266 from 150 pN to around 700 pN (Figure S2b) which was higher than adhesion forces on cell surfaces. This 267 difference could be explained by a higher density of saccharides on the model surfaces than on the cell 268 wall surface and/or the absence of interferences with other molecules present on the bacteria surface. 269 Rupture lengths however were lower on model surfaces than when SPFS was performed on cells and 270 ranged from 50 to 100 nm (Figure S2c). Molecules grafted on model surfaces were monosaccharides

whereas saccharides in *S. aureus* cell wall are mostly in the form of residues in long glycopolymer chains like WTA [35, 39-41]. The difference in the length, and thus in elasticity, between these molecules could explain the lower rupture lengths and the different FD curve signatures observed when monosaccharidecoated model surfaces were scanned by SPFS.

275 Phage-infected S. aureus proportion in presence of a monosaccharide was correlated to the adhesion 276 frequency of phage 187 on a model surface coated with the same monosaccharide (Figure 8). In other 277 terms the higher adhesion forces between a monosaccharide (GlcNAc, GalNAc, Glc and Gal) and phage 278 187 are, the better this saccharide will be in inhibiting phage infection. This relationship between adhesion 279 to model surfaces and inhibition of infection can help predict the infectivity of phage 187 on S. aureus in 280 presence of a saccharide by screening the adhesion of the phages on a saccharide-coated model surfaces. 281 It would be relevant to test with other saccharides and phages to build a predictive model by extrapolation 282 where screening adhesion forces of a phage with saccharide coated surfaces can help predict the inhibition 283 potential of this saccharide with a specific phage.

#### 284 4. CONCLUSIONS

285 With the raise of antibiotic-resistant infections, thorough understanding of phage-host adhesion becomes 286 crucial. This work demonstrates the potential of single-particle force spectroscopy, a homologous 287 technique to single-molecule force spectroscopy and single-cell force spectroscopy, to reveal major cell 288 wall receptors for phages and to help identify best suitable phage infection inhibitors. Phage-decorated 289 atomic force microscopy tips were successfully used to screen phage adhesion on single living cells and on 290 model surfaces coated with molecules of interest. Single-particle force spectroscopy evidenced affinities 291 of phage 187 for monosaccharides presenting acetyl amino groups. It also evidenced that these 292 monosaccharides were able to reduce phage adhesion on S. aureus cells after a contact of 2h. Single-293 particle force spectroscopy combined with plaque forming unit counts showed a direct correlation 294 between phage 187 infectivity in presence of N-acetyl-D-glucosamine, N-acetyl-D-galactosamine, glucose 295 and galactose and phage adhesion forces on coated surfaces with these saccharides. This suggested that 296 the higher adhesion forces between a saccharide and phage 187 are, the better this saccharide will be in 297 inhibiting phage infection of S. aureus. Single-particle force spectroscopy can help identifying the 298 relationship between the propensity of an inhibitor to bind to phages and the efficacy of inhibition in order 299 to develop models that can predict efficacy of inhibitor only by characterizing phage-inhibitor interactions, 300 for faster inhibitor characterization, and that helps choosing the best suitable inhibitor. This study expands 301 the repertoire of atomic force microscopy-based force spectroscopy and provides insights on correlation 302 between phage adhesion and infectivity.

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308 The authors declare no competing interests.

## 309 AUTHOR CONTRIBUTIONS

B.A., M.G., C.G., N.V., I.B. and S.E.K.C designed research; B.A., M.G., N.V. and S.E.K.C performed research;

B.A., M.G., C.G., N.V., I.B. and S.E.K.C analyzed data; B.A., M.G., C.G., I.B. and S.E.K.C wrote the original

312 draft.

## 313 ELECTRONIC SUPPLEMENTARY MATERIAL (ESM)

- Supplementary material (Figure S1 and S2 showing additional adhesion maps recorded on *S. aureus* cells
- and on GlcNAc-coated model surfaces with AFM tips decorated with phage 187 and Figure S3 showing
- the impact of GlcNAc on cell wall) is available in the online version of this article.

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# 433 FIGURES





Figure 1. (a) Schematic of the interaction between the phage 187 and *S. aureus*. The phage receptorbinding proteins (RBPs) recognize sub-units of the peptidoglycan-anchored wall teichoic acids (WTA) on *S. aureus* surface. (b) PFU obtained after mixing 1 x 10<sup>2</sup> PFU of purified phage 187 with approximately 1 x 10<sup>8</sup> *S. aureus* cells in TSB medium supplemented with 1 mM CaCl<sub>2</sub> and 1 mM MgSO<sub>4</sub>. A uniform bacterial mat covered the TSA agar in the Petri dish. Bacterial infections lead to lysis plaques (clear dots) of ~1 mm diameter where bacteria were lysed and did not grow. For eyes guidance, three lysis plaques are pointed with white arrows and a three-time zoom of a small area is added.

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Figure 2. (a) AFM height images in air of purified phage 187 grafted by NHS/EDC coupling on gold-coated surfaces. (b) High resolution topography tridimensional image (500 x 500 nm) in air of a single phage 187 covalently immobilized on gold substrate and corresponding cross section taken along the dashed line on the height image. (c) AFM height images in buffer of purified phage 187 grafted by NHS/EDC coupling on gold-coated surfaces. (d) Principle of single particle force spectroscopy where AFM tips decorated with phage 187 are used to probe single *S. aureus* cells.

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Figure 3. (a) Adhesion maps (500 x 500 nm<sup>2</sup>) recorded on *S. aureus* cells (insets, scale bar = 250 nm) with AFM tips decorated with phage 187. White/grey pixels correspond to adhesive events and black pixels correspond to non-adhesive events (scale 0-1000 pN). (b) Corresponding adhesion force and (c) rupture length histograms together with representative force curves. Data were obtained from three independent experiments using different tips and independent cultures.



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Figure 4. Infectivity of phage 187 towards *S. aureus* cells after 2 hours contact with free monosaccharides (10 mM) determined by plaque forming unit (PFU) counting. *S. aureus* cells (1 x 10<sup>8</sup>) were mixed with phage 187 (1 x 10<sup>2</sup>) and plated on growth media. The percentage of infection was determined by comparison to the number of PFU in absence of free monosaccharides. (Glc: Glucose; Gal: Galactose; GalNAc: N-acetyl-D-galactosamine; GlcNAc: N-acetyl-D-glucosamine; LOD: Limit of detection). Error bars correspond to standard errors calculated from three independent experiments.



Figure 5. (a) Topography images of *S. aureus* cells trapped in porous membranes (scale bar = 250 nm) and
probed with AFM tips decorated with phage 187. (b, c) Adhesion force histograms and adhesion force
maps (500 x 500 nm<sup>2</sup>, insets) obtained on cells (b) in the absence and (c) in the presence of free
monosaccharides (10 mM). White/grey pixels correspond to adhesive events and black pixels correspond

- 475 to non-adhesive events (scale 0-1000 pN). The added free monosaccharides are indicated above the
- 476 adhesion force histograms in (c).





478 Figure 6. Adhesion frequency recorded between the surface of S. aureus cells and AFM tips decorated with 479 phage 187 with or without free monosaccharides (10 mM). For each monosaccharide, adhesion maps were 480 recorded on four cells (1-4) in absence (colored bars) or presence (patterned/light colored bars) of free 481 monosaccharides (10 mM) (Purple: Glc; Blue: Gal; Green: GalNAc; Red: GlcNAc). The average decrease is 482 indicated below each monosaccharide name (mean ± s.e.). The adhesion frequency was calculated from 483 the adhesive FD curves obtained among the 1024 FD curves recorded for each map. For each 484 monosaccharide, data were obtained from three independent experiments using different tips and 485 independent cultures.



488 Figure 7. (a) AFM tips decorated with phage 187 were used to probe model surfaces coated 489 monosaccharides. (b) Topography (AFM height image; 5 x 5  $\mu$ m<sup>2</sup>; z scale = 6 nm) of GlcNAc coated 490 substrates revealing a smooth and homogeneous surface. A vertical cross section taken along the dashed 491 line is shown below the image. The thickness (~1.3 nm) was determined by imaging a square area of 1 x 1 492  $\mu$ m<sup>2</sup> at large forces (>10 nN) before imaging 5 x 5  $\mu$ m<sup>2</sup> of the same area at small forces. No scratches were 493 observed for uncoated control surfaces. (c) Boxplot of adhesion frequencies of phage 187-decorated AFM 494 tips on surfaces coated with monosaccharides. Boxes correspond to 25-75 percentiles, dashed lines to 495 means and whiskers to standard deviations. Data were obtained from two independent experiments using 496 different tips and surfaces.



Figure 8. Correlation between the percentage of infected cells in presence of different free
monosaccharides and the adhesion frequency of phages towards surfaces coated with monosaccharides
(mean ± s.e.). The red line corresponds to a linear fit for eye guidance.