# Surface-Exposed Histone-Like Protein A Modulates Adherence of *Streptococcus gallolyticus* to Colon Adenocarcinoma Cells<sup>⊽</sup>

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*Streptococcus gallolyticus* (formerly known as *Streptococcus bovis* biotype I) is a low-grade opportunistic pathogen which is considered to be associated with colon cancer. It is thought that colon polyps or tumors are the main portal of entry for this bacterium and that heparan sulfate proteoglycans (HSPGs) at the colon tumor cell surface are involved in bacterial adherence during the first stages of infection. In this study, we have shown that the histone-like protein A (HIpA) of *S. gallolyticus* is a genuine anchorless bacterial surface protein that binds to lipoteichoic acid (LTA) of the gram-positive cell wall in a growth phase-dependent manner. In addition, HIpA was shown to be one of the major heparin-binding proteins of *S. gallolyticus* able to bind to the HSPG-expressing colon tumor cell lines HCT116 and HT-29. Strikingly, although wild-type levels of HIpA appeared to contribute to adherence, coating of additional HIpA at the bacterial surface resulted in reduced binding site in HIpA. Altogether, this study implies that HIpA serves as a fine-tuning factor for bacterial adherence.

The human gastrointestinal tract is the habitat for a large and dynamic bacterial community, which is essential for intestinal epithelial homeostasis and human health. In contrast, gut flora may also be of critical importance in gastrointestinal diseases, such as colon cancer (10). Although hundreds of microbial species reside in the human intestinal tract, only a systemic infection with the gram-positive gut bacterium Streptococcus gallolyticus (formerly known as Streptococcus bovis biotype I) has a well-known association with colon cancer (13, 20, 44). (S. gallolyticus has the highest association with colorectal cancer of all S. bovis biotypes. Unfortunately, not all studies have distinguished S. bovis biotypes. Therefore, we use the name S. bovis when the specific biotype is not known and S. gallolyticus only when it is certain that the authors refer to S. bovis biotype I.) This bacterium, which can normally be detected in the gastrointestinal tracts of about 10% of the human population (31), is considered to be a low-grade opportunistic pathogen that can establish infections only in individuals with damaged heart valves (endocarditis) or a compromised immune system (bacteremia). Interestingly, fecal carriage of S. bovis was shown to be increased about fivefold in patients with colon cancer, indicating that colon tumors constitute a prefer-

\* Corresponding author. Mailing address: Department of Laboratory Medicine, Clinical Chemistry, 441, Radboud University Nijmegen Medical Centre, P.O. Box 9101, 6500 HB Nijmegen, The Netherlands. Phone: 31-24-3618947. Fax: 31-24-3541743. E-mail: H.tjalsma@akc .umcn.nl. ential colonization niche for this bacterium (20). In addition, multiple studies have shown that in up to 60% of patients with *S. bovis* endocarditis or bacteremia a colon tumor was detected upon full bowel examination (44). This strongly suggests that a colon polyp or tumor is the main portal of entry for *S. gallolyticus*, which is underscored by the fact that patients with *S. bovis* endocarditis are significantly older than those with endocarditis due to other streptococci (15). Noticeably, this could mirror the increased frequency of tumors in the elderly population.

A first requirement to establish a bacterial infection is a dependable connection between bacterial adhesins and host surface structures to withstand microbial competition and mechanical cleansing processes within the intestinal tract (14, 39). It has been suggested that heparan sulfate proteoglycans (HSPGs) present on intestinal epithelial cells are a target for S. bovis adherence and internalization, thereby participating in bacterial translocation across the intestinal epithelial barrier (12). Interestingly, the heparin-binding histone-like protein A (HlpA) from S. bovis was recently shown to be present in cell wall extracts and was a target of the humoral immune response in patients with colon cancer (38). Several other studies confirmed that this conserved bacterial protein from Streptococcus pyogenes, which is 95.6% identical to S. gallolyticus HlpA, had affinity for lipoteichoic acid (LTA), an intrinsic component of the gram-positive cell wall (22, 35, 43). Furthermore, proteomic analysis of the bacterial surface of S. pyogenes revealed that HlpA is a surface-linked protein (33). Taken together, these observations suggested that HlpA from S. gallolyticus is a

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so-called anchorless surface protein, which is a target of the humoral immune system upon infection and may mediate bacterial adherence to colon tumor cells by linking bacterial LTA to HSPG on colon epithelial cells.

Anchorless proteins comprise typical cytoplasmic proteins that are released by unknown mechanisms and reassociate with components of the bacterial cell wall or proteins at the bacterial surface (23, 36). To several of these anchorless surface proteins a secondary extracytoplasmic function has been assigned. For instance, alpha-enolase and glyceraldehyde 3-phosphate of group A streptococci have the ability to bind to plasminogen and fibronectin on pharyngeal epithelial cells (2, 24–26). Similarly, these surface-exposed proteins from *Streptococcus pneumoniae* bind to and activate plasminogen and are important for tissue invasion and virulence (5).

Therefore, the main aim of this study was to investigate the ability of *S. gallolyticus* HlpA to mediate bacterial adherence to colon tumor cells. Our experiments showed that purified HlpA did bind to both intact bacteria and colon tumor cells. Furthermore, antibody-mediated blockage of HlpA reduced the adhering properties of *S. gallolyticus*, whereas loading *S. gallolyticus* cells with purified HlpA had no promoting effect on bacterial adherence.

### MATERIALS AND METHODS

Bacterial strains and media. The strain used in this study was *S. gallolyticus* subsp. *gallolyticus* UCN34 (entitled herein *S. gallolyticus*), which was previously isolated from a colon cancer patient with coincidental endocarditis (P. Glaser, unpublished data). This strain relates to *S. bovis* biotype I, which has the highest association with colon cancer (6). *S. gallolyticus* cells were cultured at  $37^{\circ}$ C/5% CO<sub>2</sub> in brain heart infusion broth (Difco Laboratories) supplemented with 1% glucose. *S. gallolyticus* lysates were obtained by mechanical disruption of frozen bacterial cell pellets at 2,000 rpm in liquid N<sub>2</sub>. Lysed bacteria were dissolved in phosphate-buffered saline (PBS) and stored at  $-20^{\circ}$ C. The *Escherichia coli* strains DH5 $\alpha$  (Invitrogen), used for recombinant DNA procedures, and BL21(DE3) (Novagen), used as an overproduction host, were grown aerobically at 37°C in Luria broth. When required, the medium for *E. coli* was supplemented with 50 µg/ml ampicillin.

**Colon adenocarcinoma cell lines and media.** The colon adenocarcinoma cell lines Caco-2, HCT116, and HT-29 (obtained from the American Type Culture Collection) were grown in Dulbecco's modified Eagle's medium (DMEM) (Lonza) supplemented with 10% fetal calf serum, 20 mM HEPES, 100 nM nonessential amino acids, and 2 mM L-glutamine (Gibco). HT-29 and HCT116 cells were subcultured every 3 days and Caco-2 cells every 7 days. Coculture experiments were performed in six-well plates with complete DMEM containing 1% fetal calf serum.

**DNA techniques.** Procedures for PCR, DNA purification, restriction, ligation, agarose gel electrophoresis, and calcium chloride transformation of *E. coli* were carried out as described by Sambrook et al. (29). Enzymes were obtained from New England Biolabs and Applied Biosystems. To construct pET11-HlpA-His, first a fragment comprising the complete open reading frame of the *hlpA* gene of *S. gallolyticus* was amplified by PCR using the primers HlpA-u (5'-ACGTCAT ATGGCTAACAAACAAGATTT AATCGC-3'), containing an NdeI cleavage site, and HlpA-r (5'-CGTAAGGATCCTTAATGGTGATGGTGATGGTGATGGTGATTT TACAGCGT CTTTAAGTGCTTTACC-3'), which inserts a hexahistidine (six-His) tag upstream of the *hlpA* stop codon and contains a BamHI cleavage site. Next, the amplified fragment was cleaved with NdeI and BamHI and ligated into the corresponding sites of pET11a (Novagen).

**Protein purification.** To overproduce HlpA-His, 500  $\mu$ M isopropyl- $\beta$ -D-thiogalactopyranoside (IPTG) was added to exponentially growing *E. coli* BL21(DE3) cells that were first introduced with pET11-HlpA-His. Cells were harvested after 4 h and lysed by three freeze-thaw cycles. Native His-tagged HlpA was purified by nickel affinity chromatography by use of a Ni-nitrilotriacetic acid mini spin kit from Qiagen.

**SDS-PAGE and Western blotting.** To monitor the affinity of HlpA-His for intact *S. gallolyticus* cells, bacteria were incubated with purified HlpA-His ( $\sim$ 30  $\mu$ M) for 1 h at 37°C. Next, bacteria were washed three times with sterile PBS and

directly incubated in Tricine sodium dodecyl sulfate (SDS) sample buffer (30) for 15 min. Bacteria and isolated protein fractions were analyzed by Tricine SDS-polyacrylamide gel electrophoresis (SDS-PAGE) (30). To detect HIpA or HIpA-His, proteins were transferred to polyvinylidene fluoride membranes (Amersham) by Western blotting (40). Blots were incubated with monoclonal antihexahistidine antibodies (Qiagen) or highly cross-reactive anti-*S. pyogenes* HIpA antibodies that were raised against recombinant *S. pyogenes* histone-like protein and purified as described previously (16). Bound antibodies were visualized with an ECL detection system (Amersham) using anti-rabbit or antimouse immunoglobulin G horseradish peroxidase conjugates (Jackson Immuno-Research).

In-cell Western analysis. To investigate the binding of HlpA-His to intact S. gallolyticus or adenocarcinoma cells, 96-well plates were coated with either adenocarcinoma cells (50,000 cells/well) or bacteria (1  $\times$  10<sup>9</sup>/ml). Briefly, bacteria were cultured to the exponential (optical density at 600 nm of 0.6) or stationary (optical density at 600 nm of 1.0) growth phase and subsequently incubated with purified HlpA (~30  $\mu$ M) at 37°C/5% CO<sub>2</sub> for 1 h. Next, bacteria were allowed to attach to a poly-lysine-coated 96-well plate at 4°C overnight. To determine the net binding of endogenous and recombinant HlpA, coated bacteria were stripped with trypsin-EDTA (20 U/ml). The HT-29, Caco-2, and HCT116 cells were cultured in 96-well plates for 3 days at 37°C to create a confluent monolayer. Purified HlpA-His (~30 µM) was dissolved in specified growth medium and allowed to bind to confluent monolayers for 1 h. Excess protein was removed by three washes with PBS. Then, cells were fixed with 3.7% formaldehyde and left intact to determine the amount of extracellularly adhered HlpA. After fixation, plates were washed with PBS and blocked with Li-COR Odyssey blocking buffer (Li-COR Biosciences). HlpA was detected with a polyclonal HlpA antibody and visualized with Odyssey (Li-COR Biosciences) and a secondary anti-rabbit antibody conjugated with Alexa Fluor 680 (Molecular Probes).

LTA and heparin binding assays. Streptococcal LTA (1  $\mu g/\mu$ ]; Sigma) was immobilized on Interaction Discovery Mapping (IDM) affinity beads (Ciphergen Biosystems) by incubation in 100 mM ammonium acetate (pH 5). Beads were washed extensively with PBS after blocking of unoccupied binding sites by 50 mM Tris-HCl, pH 8. Immobilized LTA or heparin (heparin Sepharose; Pierce) was incubated with soluble bacterial proteins for 1 h in PBS, followed by centrifugation, after which the supernatant (unbound fraction) was collected. Next, LTA or heparin beads were washed three times with PBS, and retained proteins were eluted in 100 mM ammonium acetate (pH 6) containing 0.1% SDS at 95°C for 10 min. Protein profiles of bound and unbound low-molecular-mass proteins were generated by surface-enhanced laser desorption-ionization time-of-flight mass spectrometry, as described previously (38). Alternatively, eluted proteins were boiled in SDS sample buffer for SDS-PAGE and Western blot analysis. Protein identification was performed by in-gel tryptic digestion followed by peptide sequencing by tandem mass spectrometry, as described previously (37).

**Bacterial adherence assay.** The influence of HlpA on adherence of *S. gallolyticus* to adenocarcinoma cells (HCT116 and HT-29) was studied with an adherence assay as described previously (3). First, *S. gallolyticus* cells were treated with recombinant HlpA-His (~30  $\mu$ M), HlpA antibodies (1:1,000), or HlpA antibodies preblocked with HlpA-His in PBS. Nontreated *S. gallolyticus* cells were used to determine normal adherence levels. After pretreatment of the bacteria, the bacteria were added at a multiplicity of infection of 20:1 (bacteria/cells) to subconfluent monolayers, and the infected cells were incubated for 2 h. To determine the number of adhered bacteria, the monolayers were washed three times with PBS and lysed with ice-cold PBS containing 0.025% Triton X-100. Serial dilutions of cell lysates were plated on blood-agar plates and incubated at 37°C/5% CO<sub>2</sub> to count the CFU.

#### RESULTS

HlpA of *S. gallolyticus* is an anchorless bacterial surface protein with affinity for colon tumor cells. To act as adherence factors, anchorless proteins should have affinity for components of both bacterial and colon tumor cells. To evaluate this for HlpA from *S. gallolyticus*, the *hlpA* gene was first cloned in such a way that a hexahistidine extension (His tag) was added to the carboxyl terminus of HlpA. The resulting HlpA-His protein was overproduced and purified from *E. coli* cells to determine affinity for the surfaces of intact *S. gallolyticus* cells (Fig. 1).

To investigate whether the surface expression of HlpA from



FIG. 1. Binding of HlpA to intact bacteria. Western blot analysis of *S. gallolyticus* cells that were incubated with (+) or without (-) purified HlpA-His. In the left panel, the recombinant and endogenous HlpA molecules were visualized with anti-HlpA ( $\alpha$ HlpA) antibodies. The upper band in the "+" lane corresponds to the recombinant HlpA-His molecule that is also detected by the anti-His ( $\alpha$ His) antibody (right panel).

S. gallolyticus is also growth phase dependent, S. gallolyticus cells were harvested during the exponential and stationary growth phases and immobilized in microtiter plates for in-cell Western analysis using HlpA antibodies. To reveal the endogenous expression of HlpA at the bacterial surface, bacterial cells were stripped with trypsin-EDTA to remove extracellular proteins. This showed that the endogenous expression of surface HlpA was indeed more abundant during the stationary growth phase (P = 0.0095) than during the exponential phase (not significant) (Fig. 2), but it should be noted that some low level of autolysis could have contributed to the amount of endogenous surface HlpA. Interestingly, the extracellular addition of HlpA-His to intact bacteria showed that the cell wall of S. gallolyticus cells from the stationary growth phase was not saturated with HlpA (P = 0.0331), whereas no additional binding of HlpA-His to exponentially growing cells was observed. These data show that endogenous HlpA is expressed at the cell surface of S. gallolyticus and that this phenomenon seems present most prominently during the stationary growth phase.

To identify whether HlpA can play a role in S. gallolyticus



FIG. 3. Binding of HlpA to intact adenocarcinoma cells. In-cell Western analysis of HT-29 and HCT116 colorectal cancer cells incubated with purified HlpA-His from *S. gallolyticus*. Detection of HlpA-His was established by incubation with rabbit anti-HlpA antibody and with Alexa Fluor 680-labeled secondary anti-rabbit antibody. The bars represent the fluorescence intensities of HlpA-His on adenocarcinoma cells. The fluorescence intensity of HlpA was corrected for the basal fluorescence level (nonspecific fluorescence) of the indicated cell line. The blank measurement was obtained by subtracting the fluorescence intensity of a blank well incubated with DMEM from that of a well incubated with DMEM with HlpA (background measurement). Error bars indicate the standard errors from two replicate experiments.

adherence, the binding of HlpA-His to the adenocarcinoma cell lines HT-29, HCT116, and Caco-2 was quantified by in-cell Western analysis. The fluorescence intensity of HlpA was highest in HT-29 cells and then in HCT116 cells compared to that for the blank measurement (Fig. 3). In contrast, the fluorescence intensity of Caco-2 adenocarcinoma cells after incubation with HlpA-His remained similar to that for the blank measurement, which may reflect the low level of HSPGs in these cells (reference 11 and data not shown). Taken together, these data show that extracellularly added HlpA can bind to



FIG. 2. Affinity of HlpA for *S. gallolyticus* in exponential and stationary growth phases. In-cell Western analysis of *S. gallolyticus* bacteria in exponential and stationary growth phases exposed to HlpA-His. Approximately  $1 \times 10^9$  exponential- or stationary-phase-grown bacteria were coated onto a 96-well plate. Detection of HlpA and HlpA-His was established by rabbit anti-HlpA antibodies and secondary Alexa Fluor 680-labeled anti-rabbit antibody. The bars represent the fluorescence intensities of extracytoplasmic recombinant HlpA-His and endogenous HlpA of bacteria. Note that nonstripped (top row, -) bacteria have in addition to HlpA-His, endogenous HlpA at their surfaces, whereas trypsin-stripped (top row, +) or without (bottom row, -) HlpA-His. The fluorescence intensity was corrected for background measurement of the incubation medium. Error bars indicate the standard errors from two replicate experiments. \*, P < 0.05; \*\*, P < 0.01; ns, not significant (Student's *t* test).



the bacterial surface and also has affinity for HT-29 and HCT116 colon tumor cells and thus, in principle, should be able to act as a mediator for bacterial adherence.

HlpA from S. gallolyticus is a heparin- and LTA-binding protein. To gain evidence for the components on the colon tumor and bacterial cell surfaces that are responsible for the retention of HlpA, in vitro binding assays were performed. Previous studies showed that HlpA from S. bovis biotype II is a heparin-binding protein (37) and indicated that HSPGs on epithelial cells may have a role in the binding and invasion of these bacteria (12). To confirm heparin affinity of endogenous HlpA from the S. gallolyticus strain used in this study and that of its recombinant analogue HlpA-His, cell lysates and purified HlpA-His were incubated with immobilized heparin. As shown in Fig. 4A, both endogenous and recombinant HlpA(-His) had a high affinity for heparin. Furthermore, mass spectrometry analysis of the low-molecular-mass proteins showed that heparin affinity is selective for HlpA (Fig. 4B and C), which appeared to be the major low-molecular-mass heparin-binding protein of S. gallolyticus. To investigate whether S. gallolyticus contains additional heparin-binding proteins of >15 kDa, S. gallolyticus total cell lysates were incubated with immobilized heparin, after which retained proteins were analyzed by SDS-PAGE. These analyses confirmed that HlpA is one of the major heparin-binding proteins from S. gallolyticus; however, at least seven additional heparin-binding proteins were observed (Fig. 5A), and of these, protein 2 was identified as a putative alpha/beta hydrolase of 44 kDa (Fig. 5B). Nevertheless, these additional proteins were not present or scarcely present in S. gallolyticus cell wall extracts (our unpublished observations). Thus, HlpA appears to be the major surfaceexposed heparin-binding protein from S. gallolyticus.

To show LTA affinity of endogenous and recombinant HlpA(-His) from *S. gallolyticus*, cell lysates and purified HlpA-His were incubated with immobilized LTA and retained proteins were analyzed by mass spectrometry. As shown in Fig. 4B and C, endogenous and recombinant HlpA(-His) both had affinity for LTA. This suggests that HlpA is retained by LTA in the bacterial cell wall and can mediate bacterial adherence by linking HSPGs of epithelial cells to LTA of the bacterial cell wall. Importantly, these analyses also indicated that the native features of HlpA have been preserved in recombinant HlpA-His, which thus can be used as an experimental substitute for endogenous HlpA.

Surface HIpA affects *S. gallolyticus* adherence to colon tumor cells. To investigate whether surface HIpA indeed mediates the adherence of *S. gallolyticus* to colorectal cancer cells, bacterial adherence assays were performed. First, endogenous HIpA on the surface of *S. gallolyticus* was blocked by the anti-HIpA immunoglobulin G that was also used for in-cell

Western analysis (Fig. 2 and 3). As shown in Fig. 6A, this resulted in approximately 20% (not significant) and 60% (P = 0.0158) reduced adherence of *S. gallolyticus* to HCT116 and HT-29 cells, respectively, compared to the level for untreated *S. gallolyticus* cells (100% adhesion). The latter inhibition was partially restored (not significant) by antibodies preblocked with HlpA-His that were used to confirm a specific interaction between HlpA and the anti-HlpA antibody.

To investigate whether additional surface HlpA could further increase bacterial adherence, *S. gallolyticus* cells were coated with HlpA-His prior to use in an adherence assay. Unexpectedly, adhesion of HlpA-His-coated bacteria did not result in an increase, but decreased the adherence to HT-29 (not significant) and HCT116 (P = 0.0242) cells to about 65% of the level for untreated *S. gallolyticus* cells (Fig. 6B). This implies that an excess of surface HlpA has an inhibitory effect.

Heparin and LTA compete for the same binding sites in HlpA. The above-described apparent contradictory data suggest that HlpA adds to bacterial adherence but point to a more complex process than simply linking HSPGs and LTA. One of the influencing factors could be that HSPGs and LTA, both strong negative structures, compete for the same positively charged binding sites in HlpA (28, 35). Therefore, the interference of LTA in heparin binding of HlpA was investigated by mass spectrometry. As shown in Fig. 7, preincubation of HlpA with soluble LTA inhibited the binding of both HlpA and HlpA-His to immobilized heparin in a concentration-dependent manner. In contrast, incubation with LTA after binding to immobilized heparin had no significant inhibitory effect on the binding of HlpA and HlpA-His to heparin. The fact that these effects were more pronounced with HlpA-His is possibly due to the scavenging of LTA by other S. gallolyticus proteins with affinity for LTA (and possibly heparin) in the total cell extracts (Fig. 4B). Thus, these data suggest that HlpA cannot efficiently bind to both HSPG and LTA simultaneously.

## DISCUSSION

HSPGs are proposed to give bacteria the opportunity to adhere to and invade the epithelium, but whether bacteria can use these eukaryotic structures is dependent on bacterial heparin-binding proteins at the bacterial surface. Several studies have stated that histone-like proteins are detected in the culture supernatant or at the bacterial cell surface of *Helicobacter pylori*, *S. pyogenes*, and *Streptococcus intermedius* (19, 21, 33). Furthermore, it has been shown previously that HlpA from *S. pyogenes* can be complexed to LTA (33, 35). Therefore, we hypothesized that the histone-like protein HlpA of *S. gallolyticus* was a likely candidate to mediate adherence via these structures. Our studies clearly showed that endogenous and

FIG. 4. Heparin and LTA affinity of HlpA. (A) Western blot analysis of a total *S. gallolyticus* (Sga) cell lysate (CL) and purified (P) HlpA-His and the respective fractions that remain unbound (U) after incubation with immobilized heparin and that have heparin affinity (H). Blots were decorated with anti-HlpA or anti-six-His antibodies. The positions of endogenous HlpA and recombinant HlpA-His are indicated. (B) Low-molecular-mass protein profiles of a total *S. gallolyticus* cell extract, the respective proteins that remain unbound after incubation with immobilized heparin or LTA, and the respective proteins that have heparin or LTA affinity. Note that a second (unknown) protein with both heparin and LTA affinity is indicated with an asterisk. (C) Low-molecular-mass spectra of purified HlpA-His from *S. gallolyticus*, the fraction that remains unbound after incubation with immobilized heparin or LTA, and the proteins that have heparin or LTA affinity. The 9,707- and 10,530-Da peaks, corresponding to, respectively, endogenous HlpA and recombinant HlpA-His, are indicated. The peak intensity is given in arbitrary units.



FIG. 5. (A) SDS-PAGE analysis of a total *S. gallolyticus* cell lysate (CL) and the respective proteins that remain unbound (U) after incubation with immobilized heparin or have heparin affinity (H). The position of endogenous HlpA is indicated. Unknown heparin-binding proteins 1 to 7 are indicated with asterisks. M, molecular mass marker. (B) Identification of protein band 2 as a putative alpha/beta hydrolase of *S. gallolyticus*.

with immobilized heparin or have heparin affinity (H). The position of endogenous HIpA is indicated. Unknown heparin-binding proteins 1 to 7 are indicated with asterisks. M, molecular mass marker. (B) Identification of protein band 2 as a putative alpha/beta hydrolase of *S. gallolyticus*. Band 2, a putative heparin-binding protein (indicated in panel A), was excised from the gel and digested with trypsin to allow tandem mass spectrometry and peptide mass fingerprinting. The tandem mass spectrometry spectrum of peptide "QKYLINSILKW," m/z 1,405.82, is shown. The theoretical series of b ions produced from cleavage of the amide bond and y ions produced by cleavage of the amide bond are indicated, with the actual identified ions printed in bold. The theoretical molecular mass of the corresponding putative alpha/beta hydrolase is 43.7 kDa, which is in-line with its mobility by SDS-PAGE (see panel A).



FIG. 6. Surface HlpA modulates bacterial adherence. (A) *S. gallolyticus* cells were incubated with (+) or without (-) anti-HlpA ( $\alpha$ -HlpA) antibodies. Antibodies preincubated with HlpA-His (HlpA- $\alpha$ -HlpA) served as a control. Adherence of *S. gallolyticus* (Sga) to HCT116 and HT-29 cells was monitored by CFU serial dilution counting. The adherence of *S. gallolyticus* for the specified cell lines was set at 100%. Error bars indicate the standard errors from three replicate experiments. \*, P < 0.05 (Student's *t* test). (B) Adherence of *S. gallolyticus* to the specified cell serial dilution counting. The adherence of by CFU serial dilution counting to the specified cell series and the standard errors from three replicate experiments. \*, P < 0.05 (Student's *t* test). (B) Adherence of *S. gallolyticus* to the specified cell lines was set at 100%. Error bars indicate the standard errors from three replicate the standard errors from three replicate the standard errors indicate the standard errors have set at 100%. Error bars indicate the standard errors from three replicate experiments. \*, P < 0.05 (Student's *t* test).

recombinant HlpA(-His) have affinity for both heparin and LTA. Our data demonstrate that endogenous HlpA is preferentially expressed at the bacterial surface in the stationary growth phase, which is in line with the study of Katsube et al., who reported that the histone-like protein MDP1 of Mycobacterium smegmatis accumulates in cell wall fractions in the stationary phase (18). Furthermore, our findings correlate with previous studies reporting the release of HlpA during the stationary growth phase (22, 35). In the exponential phase, HlpA is required intracellularly to execute its physiological role in nucleoid formation (1), but even though endogenous HlpA is maintained intracellularly, recombinant HlpA-His was not able to bind to the bacterial surface. Thus, HlpA relocation at the bacterial surface is growth phase dependent in vitro, which could mirror the in vivo bacterial steady-state situation in, for instance, the gastrointestinal tract.

A previous study by Henry-Stanley et al. showed that the colon adenocarcinoma cell line HT-29 expresses HSPGs and



FIG. 7. LTA inhibition of heparin binding of HlpA. Low-molecular-mass profiles of heparin-binding proteins from a total *S. gallolyticus* cell extract or that of purified HlpA-His were generated by mass spectrometry. Bars represent the peak intensities of HlpA (m/z 9,707) and HlpA-His (m/z 10,530) that were retained by immobilized heparin after preincubation or postincubation with 25 or 50 µg/µl soluble LTA. Error bars indicate the standard errors from three replicate experiments.

more specifically the heparan sulfate syndecan-1, whereas Caco-2 cells have only low levels of these proteoglycans (11). Furthermore, Stinson et al. reported that bacterial HlpA of S. pyogenes binds to heparan sulfate in the extracellular matrix of human epithelial Hep-2 cells (35). We observed that recombinant HlpA-His could bind to HT-29 and HCT116 adenocarcinoma cells but not to Caco-2 cells, which coincides with the amount of HSPGs and syndecan-1 and with the previously reported conclusion that S. bovis may use heparan sulfates to invade eukaryotic cells (12). Furthermore, we found that S. gallolyticus adherence to both HCT116 and HT-29 cells was reduced by anti-HlpA antibodies, although the effects seen in the independent experiments did not reach statistical significance. Nevertheless, the same trend was observed for both cell lines, which is in line with the fact that heparin inhibited S. bovis internalization to a similar extent in HT-29 cells (12). Unexpectedly, however, when S. gallolyticus was coated with additional recombinant HlpA-His, adherence was decreased. This implies that an excess of surface HlpA blocks adherence. A possible explanation for these apparent contradictory effects may be given by the fact that LTA competes with heparin for the binding of endogenous and recombinant HlpA, indicating that HSPGs and LTA use the same binding site(s) in this molecule (Fig. 7). The E. coli DNA binding protein HU, which is homologous to HlpA, has been studied extensively in its interaction with DNA. Homotypic dimers of E. coli HU interact with DNA via its binding arms, which resemble the amino acid sequence of S. gallolyticus HlpA (27). These binding arms display an amino acid sequence that contains a typical positively charged heparin-binding site, as proposed by Cardin and Weintraub (4, 9), which interacts with negatively charged sulfate or carboxyl groups on heparin chains but may also interact with the negatively charged LTA (42).

Thus, although we have shown that HlpA is a major heparinbinding protein at the bacterial surface, the exact contribution of HlpA and HSPGs in the adherence of S. gallolyticus to colon tumor cells is not yet fully resolved. In this respect, it was interesting to note that Sillanpaa and coworkers showed that S. gallolyticus can bind to several eukaryotic extracellular matrix proteins (34). Others demonstrated that LTA itself is important for bacterial adherence and invasion (7, 17, 32); this was confirmed for S. gallolyticus in a study by Von Hunolstein et al., who showed that adherence to buccal epithelial cells was affected by anti-LTA antibodies and epithelial LTA treatment (41). In view of our current observations, increasing the amount of HlpA at the bacterial surface could therefore decrease the amount of interacting LTA molecules, when attachment of our clinical isolate is mainly LTA driven. Furthermore, our results could imply that excess surface HlpA hinders the interaction of other important adhesion molecules on S. gallolyticus.

For opportunistic pathogens, such as S. gallolyticus, a temporal regulation of interactions with host cells may be of critical importance. Although a firm connection with host epithelial (tumor) cells is key during the first phase of infection, an interaction with tissue macrophages that form the second line of defense against invading pathogens should be avoided. In this respect, surface HlpA may for instance shield LTA from interacting with Toll-like receptors on macrophages (8). Our current data show that surface HlpA may be one of the modulators of these bacterial interactions with host cells. However, HlpA from Streptococcus mitis by itself has been shown to activate an immune response in murine macrophages in vitro, which is mediated by the induction of interleukin-1 and tumor necrosis factor alpha (43). Similarly, Liu et al. have shown that HlpA of S. intermedius can induce cytokine production in human macrophages (22). However, none of these studies has determined the net effect of the HlpA-mediated shielding of LTA from intact bacteria on the interaction with, and activation of, macrophages; this will be a subject of our future investigations.

Taken together, our current data underscore that bacterial adherence is a highly balanced process involving multiple interactions on the host-pathogen interface. Importantly, this study presents the first evidence that surface HIpA may provide opportunistic pathogens, such as *S. gallolyticus*, with the ability to adjust their host-pathogen interactions during different stages of infection.

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