

Pneumococcal neuraminidase A: an essential upper airway colonization factor for *Streptococcus pneumoniae*

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SUMMARY

Streptococcus pneumoniae colonizes the upper respiratory tract from where the organisms may disseminate systemically to cause life threatening infections. The mechanisms by which pneumococci colonize epithelia are not understood, but neuraminidase A (NanA) has a major role in promoting growth and survival in the upper respiratory tract. In this article we show that mutants of S. pneumoniae D39 deficient in NanA or neuraminidase B (NanB) are abrogated in adherence to three epithelial cell lines, and to primary nasopharyngeal cells. Adherence levels were partly restored by nanA complementation in trans. Enzymic activity of NanA was shown to be necessary for pneumococcal adherence to epithelial cells, and adherence of the nanA mutant was restored to wild-type level by pre-incubation of epithelial cells with Lactococcus lactis cells expressing NanA. Pneumococcal nanA or nanB mutants were deficient in biofilm formation, while expression of NanA on L. lactis or Streptococcus gordonii promoted biofilm formation by these heterologous host organisms. The results suggest that NanA is an enzymic factor mediating adherence to epithelial cells by decrypting receptors for adhesion, and functions at least in part as an adhesin in biofilm formation. Neuraminidase A thus appears to play multiple temporal roles in pneumococcal infection, from adherence to host tissues, colonization, and community development, to systemic spread and crossing of the blood-brain barrier.

INTRODUCTION

Streptococcus pneumoniae is a major human pathogen associated with a wide range of disease conditions including otitis media, sinusitis, pneumonia and meningitis (Mitchell, 2003). Pneumococci frequently colonize the nasopharynx (Watt et al., 2004; King, 2010), and from there they may disseminate to the lungs, or haematogenously, and cause disease. The pneumococcus produces a number of well-characterized virulence factors important in pathogenesis of infections (Bergmann & Hammerschmidt, 2006; Löfling et al., 2011). These include the capsule (Cps), pore-forming toxin pneumolysin (Ply), surface proteins PsaA, PspA, PspC, PhtD, PavA, and PsrP (Paterson & Orihuela, 2010), hyaluronidase, and neuraminidases (sialidases) (Mitchell, 2003). Colonization is intimately related to the mechanisms of pathogenesis of disease, but rela-

tively less is understood about the processes of initial establishment and subsequent microbial community development. In particular, the mechanisms by which pneumococci adhere to epithelial cells are relatively uncharted, and are suggested to include recognition of surface bound matrix proteins, such as fibronectin (van der Flier *et al.*, 1995 Holmes *et al.*, 2001), and glycoconjugate receptors (King, 2010; Limoli *et al.*, 2011).

Streptococcus pneumoniae produces at least three sialidases (Xu et al., 2011) with NanA and NanB proteins being the most well-characterized, and NanC present in only about 50% of strains (Pettigrew et al., 2006). Both NanA and NanB are secreted proteins (Berry et al., 1999) but only NanA carries a carboxyterminal anchorage motif to secure it to the cell wall (Camara et al., 1994). The two enzymes have different pH optima (NanA pH 6.5, NanB pH 4.5), and while NanA catalyzes the removal of Neu5Ac from a range of substrates, NanB is an intramolecular trans-sialidase selective for a2,3-sialosides (Xu et al., 2011). The sialidases each have a carbohydrate-binding module in addition to the catalytic domain that cleaves sialoglycosides to release N-acetylneuraminic acid (Neu5Ac or NANA) (Uchiyama et al., 2009).

Neuraminidase A is most highly expressed at the transcriptional level and is present in all strains (King et al., 2005). Production of NanA is up-regulated by interaction of pneumococci with host cells (LeMessurier et al., 2006) and by free sialic acid (Yesilkawa et al., 2008). In a number of studies, NanA has been shown to contribute to nasopharyngeal colonization, spread of pneumococci to the lungs (Orihuela et al., 2004), sepsis (Manco et al., 2006), development of otitis media (Tong et al., 2000), and passage across the blood-brain barrier (Uchiyama et al., 2009). By contrast, other studies have shown no effect of nanA knockout on the ability of encapsulated strain D39 to colonize mice (King et al., 2006), and no attenuation in an infant rat model of colonization (King et al., 2004). In addition, D39AnanA mutants were unaffected in adherence levels to two epithelial cell lines (King et al., 2006). However, it was also demonstrated that a $\Delta nanA$ mutant of strain R6, which is an avirulent non-encapsulated derivative of strain D39, was impaired in adherence to epithelial cells compared with parental strain R6 (King et al., 2006). There is, therefore, conflicting evidence regarding potential roles for NanA in pneumococcal adherence to mucosa and in pneumococcal colonization.

Manipulation of complex carbohydrates by pneumococci has at least two functions. Firstly, deglycosylation of host tissue receptors reveals different glycan structures for adhesins on the pneumococcal cell surface to interact with. Secondly, sequential deglycosylation of complex carbohydrates, e.g. mucins, releases sugars as carbon and energy sources for growth. In these respects, NanA, and surface protein BagA (encoding a β -galactosidase), have been shown to deglycosylate structures on host cell surfaces to reveal cryptic receptors for adherence (King et al., 2006; Limoli et al., 2011). In confirmation, S. pneumoniae cells show increased binding to sialidase-treated chinchilla trachea cells in an otitis media model (Tong et al., 2002). It is suggested that sialic acid itself has a direct role in promoting pneumococcal biofilm formation (Trappetti et al., 2009), nasopharyngeal colonization, and translocation to the lungs (Trappetti et al., 2009).

Several studies have shown specifically that NanAdeficient pneumococci are abrogated in colonizing the nasopharynx, trachea, and lungs of rodents following intranasal challenge (Tong et al., 2000; Orihuela et al., 2004; Manco et al., 2006). In addition, NanB deficient pneumococci are impaired in colonization and are unable to cause sepsis following intranasal infection (Manco et al., 2006). In this article we have further developed the hypothesis that NanA and NanB might be essential for pneumococcal colonization of nasopharyngeal and respiratory tract epithelia, processes defined in vitro by adherence and community development. In these studies we have utilized encapsulated strain D39, in contrast to many previous investigations that have employed non-encapsulated (non-pathogenic) derivatives, such as strain R6. We demonstrate that NanA activity is essential for strain D39 adherence to four epithelial cell types and that NanA behaves as a biofilm determinant. It is also shown that adherence and colonization deficiencies of a nanB mutant may result, at least in part, from decreased expression of NanA.

METHODS

Bacterial strains and growth conditions

Streptococci were grown in Todd-Hewitt broth containing 0.5% yeast extract, in sealed tubes at 37°C without shaking. *Lactococcus lactis* MG1363 was grown in M17 medium containing 0.5% glucose. Antibiotic concentrations were 5 μ g ml⁻¹ erythromycin and 10 μ g ml⁻¹ spectinomycin. *Streptococcus pneumoniae* D39 and *nanA* (Berry & Payton, 2000) and *nanB* (Berry *et al.*, 1999) mutants were kindly provided by J. Paton. Strains 25925 (serotype 23F) and 45567 (serotype 14) were kindly provided by M. Smith (HPA, Taunton, UK).

Cloning and expression of nanA gene

The nanA gene was PCR amplified with high-fidelity polymerase (Phusion, New England Biolabs) from strain D39 DNA using primers NanAF2 GGAGGATCCAAATGTCTTATTTCAGAAATCGGGA-TATA and NanAR2 GATCTGCAGTTA TTCTTGTC TCTTTTTCCCTAGCGT (BamH1 and Pstl sites in bold). The fragment was ligated into plasmid pKS80 (5.24 kb; Hartford et al., 2001), or pKS80 aad9, digested with BamH1 and Pstl, such that nanA expression was under transcriptional and translational control by endogenous vector sequences. Streptococcus gordonii was then transformed as previously described (Haisman & Jenkinson, 1991) with selection for erythromycin resistance (Em^{R} , 5 µg ml⁻¹). Plasmid pKS80 aad9 was generated as follows: the aad9 gene encoding spectinomycin resistance (Sm^R) was cut from pFW5 (Podbielski et al., 1996) with a combination of Sall/Xmal and cloned into pUC19 similarly digested with Sall/Xmal. The aad9 gene was then removed from pUC19 by digestion with Pstl, ligated into the unique Pstl site of pKS80 nanA⁺, and transformed into S. gordonii DL1 with selection for Em^R and Sm^R. Plasmids were extracted from S. gordonii and transformed into L. lactis MG1363 by electroporation as previously described (Jakubovics et al., 2005b).

Neuraminidase activity

To determine neuraminidase activity bacterial cells were grown to late exponential phase, harvested at 5000 g, washed twice in PBS and frozen at -70 °C for 16 h. Cells were then suspended in 40 mm tris-HCl (pH 7.4), disrupted by shaking with glass beads and the supernatant was removed and stored at -70 °C for 24 h. Portions of cell-free supernatants were assayed for neuraminidase activity with 2-*O*-(*o*-nitrophenyl)-a-D-*N*-acetylneuraminic acid (*o*NP-NANA)

(Manco *et al.*, 2006). Briefly, 25 μ l aliquots of cell supernatants were added in triplicate to the wells of a microtitre plate and an equal volume of 0.3 mM *o*NP-NANA in dilution buffer (100 mM citric-acid phosphate, 25 mg ml⁻¹ bovine serum albumin, 10% sodium azide; pH 6.6) was added. Samples were incubated at 37 ° for 2 h when the reaction was stopped by addition of 100 μ l ice-cold 0.5 M Na₂CO₃. Once the solution had reached room temperature the absorbance at 405 nm was read. Standard curves were constructed with clostridial neuraminidase (Sigma, Poole, UK) and known concentrations of *p*-nitrophenol.

Cell culture

A549 cells (type II pneumocytes) and HEp-2 cells (derived from human larynx carcinoma) were maintained in Dulbecco's Modified Eagle Medium (DMEM) supplemented with 10% foetal calf serum, 2 mm L-glutamine and 0.5 µg ml⁻¹ hydrocortisone. Detroit 562 cells (human pharyngeal carcinoma) were maintained in Eagle's minimum essential medium (EMEM) with 10% FCS, 1 mm sodium pyruvate, 2 mm L-glutamine, 100 U ml⁻¹ penicillin and 100 μ g ml⁻¹ streptomycin. Cultured cells were maintained in humidified air (5% CO₂ at 37 °C) and were split twice weekly or as required by trypsinization. Primary cells (nasal brushings from a healthy child) were grown in bronchial epithelial cell basal medium (BEBM) with bulletkit supplements (bovine pituitary extract, 2 ml; hydrocortisone, 0.5 ml; hEGF, 0.5 ml; epinephrine, 0.5 ml; transferrin, 0.5 ml; insulin, 0.5 ml; retinoic acid, 0.5 ml; triiodothyronine, 0.5 ml; GA-1000, 0.5 ml). Cells were grown to 80% confluence in 75 cm² culture flasks (Corning) and collected every 3-4 days (7 days for Detroit 562 cells) by washing with Dulbecco's phosphate-buffered saline (DPBS) and treating with trypsin-EDTA. Detached cells were centrifuged for 10 min (900 rpm), suspended in 5 ml appropriate growth medium and then seeded into fresh flasks using a volume equivalent to 0.5×10^6 cells ml⁻¹. Flasks were maintained in a humidified atmosphere (5% CO2 at 37 °C). All solutions were from Cambrex unless otherwise stated.

Bacterial adherence to epithelial cells

Cultured cells were seeded into 24-well plates at a density of 1×10^4 cells per well in DMEM as above

and grown to 80% confluence (48-72 h) either directly on the plastic wells, or on IMS treated glass cover slips. To encourage attachment, these cover slips had been washed in industrial methylated spirit prior to use, and in the case of primary cells, incubated for 1 h with 0.5 ml 0.3% bovine dermal colla-International). Cover gen (VWR slips were transferred to a fresh well plate that had been blocked with 1% BSA overnight and then rinsed with PBS before fresh DMEM (or BEBM for primary cells) was added. To assay bacterial adherence, early stationary phase cells of S. pneumoniae were collected by centrifugation (9000 \times g, 10 min), washed once with DMEM, and then suspended in DMEM $(OD_{600} = 0.6 \pm 0.02)$. Wells were then inoculated with 0.2 ml pneumococcal cell suspension (1×10^8) bacterial cells). Plates were routinely incubated for 45 min at 37 °C in 5% CO2. Numbers of cells adhered were determined by direct microscopic count, following Gram-stain, with an Olympus CX41 light microscope at 400× magnification. Fields of cells were selected randomly (on condition of >70% confluence and homogenous staining), and total numbers of pneumococci adhered to 500 human cells were counted. Cover slips were examined in triplicate for each treatment and adherence was expressed as numbers of pneumococci bound per 10 human cells.

In experiments utilizing the plastic wells directly, wells were rinsed with PBS and then inoculated with 1×10^8 bacterial cells (200 µl suspension $OD_{600} = 0.6$) in serum-free DMEM. Cells were incubated for up to 4 h at 37 °C in humidified 5% CO₂. The monolayers were washed with PBS, and the numbers of bacteria associated with the epithelial cells were determined as colony forming units (CFU) from dilution counts on Todd Hewitt Agar.

In some experiments, cultured cells were incubated at 37 °C with purified clostridial sialidase (Sigma), or with *L. lactis* cells, washed five times with DMEM, and then assayed for pneumococcal cell adherence by methods described above.

Biofilm formation

Biofilms were produced on saliva-pellicle coated glass cover slips following 18 h incubation at 37 °C. Biofilms were stained with FITC for microscopic imaging, or biomass quantified by crystal violet assay

as previously described (Maddocks *et al.*, 2011). Pneumococcal biofilms were grown in THY medium, while *S. gordonii* and *L. lactis* biofilms were produced by growth in modified C medium (Maddocks *et al.*, 2011).

Statistics

Results were analysed using GraphPad Prism software version 4 and are shown as mean \pm SD. Two tailed Student's *t*-test analyses were performed on data sets and differences were considered significant when P < 0.05.

RESULTS

Time course of pneumococcal adherence to epithelial cells

In initial studies we established conditions for determining adherence of S. pneumoniae to three epithelial cell lines and to primary human nasopharyngeal cells. One of the difficulties in these assays was that the production of pneumolysin affected epithelial cell integrity, a problem that we addressed previously by utilizing a pneumolysin-deficient mutant (Pracht et al., 2005). However, to address the contextual role of NanA we did not adopt this route, rather we designed assays such that the lytic effects of pneumolysin might be minimized. We measured the time course of adherence of encapsulated S. pneumoniae D39 cells, and it was clear that lysis of epithelial cells started to become apparent around 60 min after infection. In these studies the assay cultures were stained by Gram's method to reveal dark purple pneumococci on a pink background of epithelial cells, with the nuclei staining darker pink. The kinetics of interaction of S. pneumoniae D39 with A549 epithelial cells is shown in Fig. 1. There was an approximately linear increase in the numbers of bacterial cells attaching per epithelial cell that began to level off between 45 and 60 min. The numbers of pneumococci then began to increase sharply concomitant with decrease in epithelial cell integrity (Fig. 1B). By 2 h there was marked increase in pneumococcal cell numbers (Fig. 1A) and by 4 h there was evidence for extensive pneumocccal cell multiplication promoted by lytic products from the epithelial cells (Fig. 1B, panels e and f). Similar results were obtained with a Hep-2 cell



Figure 1 Time course of adherence of *Streptococcus pneumoniae* D39 to A549 cells. Panel A, numbers of *S. pneumoniae* cells attached to A549 epithelial cells (pneumocytes) were determined by microscopic count as described in Methods. Data points are means \pm SD of triplicates from three experiments. Panel B, microscopic analysis of A549 cells infected with *S. pneumoniae* D39 (Brown-Brenn stained). Numbers of bacteria (purple) adhering can be seen to increase over the first 45–60 min (panels a-d). By 2 h (panel e) cell lysis was apparent, and at 4 h (panel f) the A549 cell monolayer was almost completely destroyed.

line, and accordingly, adherence assays were performed for 45 min.

We then determined the numbers of pneumococci attached to monolayers of different epithelial cell types at approximately 10^5 epithelial cells per well. Microscopic counts of numbers of bacteria adhered to 100 epithelial cells over numerous fields of vision and triplicate wells were determined for four epithelial cell types: A459 human lung pneumocytes; Hep-2 human laryngeal carcinoma; Detroit 562 human pharyngeal carcinoma; and primary culture of naso-pharyngeal cells. Actual numbers of pneumococci wild-type D39 attached varied considerably for each cell type, with A549 cells supporting the highest levels of adherence, and Detroit 562 cells showing the lowest (Fig. 2A). There were no obvious differences

Figure 2 Adherence of *Streptococcus pneumoniae* D39 wild-type and isogenic mutants to four types of cultured epithelial cells as indicated. Panel A, numbers of adherent pneumococci were determined by microscopic count as described in Methods. Values were obtained from three independent experiments and error bars are \pm SD. **P* \leq 0.05 compared to D39. Panel B, microscopic images of adherence of *S. pneumoniae* D39 wild-type, UB1816 *nanA* or UB1817 *nanB* mutants to cultured epithelial cell types as indicated on the left sides of the panels. Adherence of pneumococci (purple) to the cultured cells (following Brown-Brenn staining) can be clearly visualized (arrows) for D39 wild-type much less so for *nanA* or *nanB* mutants (50–80% reduced levels of adherence).



in the patterns of pneumococcal cell adhesion to the epithelial cells, but there was noticeable adherence of bacteria to the regions of cytoplasmic extension for A549 and Hep-2 cells (Fig. 2B). Moreover, Detroit 562 cells were more rounded with less uniform coverage.

Neuraminidase activity is associated with adherence

To determine if enzymic activity was necessary for adherence of pneumococci to A549 cells, strains of pneumococci expressing a range of neuraminidase activities were tested for adherence in the presence or absence of the neuraminidase inhibitor Dehydro-NeuNAc (Kumar et al., 1981). The assay conditions measure principally NanA activity (Xu et al., 2011) as NanB is a trans-sialidase with pH optimum = 4.5 (Berry et al., 1996). Neuraminidase activities in cellfree extracts of strain D39 were highly sensitive to inhibition by dehydro-NeuNAc (Fig. 3A), as was commercially available clostridial sialidase (Fig. 3A). Adhesion levels of three pneumococcal strains, representing low, medium and high producers, were reduced in the presence of Dehydro-NeuNAc (Fig. 3 B), with the high-producing strain 45567 (serotype 14) being \sim 80% inhibited. These data suggested that neuraminidase activity might be a major requirement for the adherence and colonization processes.

Effects of *nanA* or *nanB* mutations on neuraminidase activity and adherence

We then examined the effects of nanA or nanB mutations on neuraminidase activity and adherence to cultured epithelial cells. Inactivation of nanA reduced neuraminidase A activity by >99% (Table 1) and led to >70% reduction in adherence levels to each of the cell cultures (Fig. 2A). Mutation of nanB led to 52% reduction in total neuraminidase activity (Table 1). It is important to note that neuraminidase B activity is optimal at pH 4.5 and therefore does not contribute significantly to the cell-associated neuraminidase activity measured at pH 7.0 in the assays utilized here. The nanB mutant strain showed 50-80% reduced adherence levels to each of the cell cultures, compared to wild-type D39 levels (Fig. 2A). The microscopic images (Fig. 2B) show clearly the effects of nanA or nanB mutations on the numbers of pneumococcal cells adhered to each of the cultured epithelial cell types.

Previous work has demonstrated that mutation in the *pavA* gene, encoding a fibronectin-binding protein (Holmes *et al.*, 2001), led to reduced adherence to epithelial cells (Pracht *et al.*, 2005), and to deficiency in colonization of the respiratory tract (Kadioglu *et al.*, 2010). The adhesion and infection phenotype of the *pavA* mutant was thus very similar to that of the *nanA* mutant. However, the *pavA* mutant UB1341



Figure 3 Inhibition of neuraminidase activity and effect on pneumococcal adherence to A549 cells. Panel A, dose responsive inhibition of neuramindase activity from *Clostridium* or *Streptococcus pneumoniae* D39 by 2,3-dehydro-2-deoxy-N-acetylneuraminic acid (dehydro-NeuN-Ac). Panel B, effect of pre-treatment of cells from low (UB979), medium (D39) or high (UB1050) level neuraminidase-producing strains of *S. pneumoniae* with 2 mM dehydro-NeuNAc on adherence levels to A549 cells. Error bars are ±SD from three independent experiments, * $P \le 0.05$.

Table 1	Bacterial	strains	utilized	in th	nis s	study	and	their	neuraminidase	activities
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Strain	Genotype	Source/Reference	Neuraminidase activity ($U \times 10^{-3}/10^{10}$ cells) ± SD	
UB 1830	S. pneumoniae D39 (wild-type) capsular serotype 2	J. Paton/laboratory stock	52.0 ± 6.30	
UB 1816	D39 nanA::ermAM	Berry & Payton (2000)	4.10 ± 1.80	
UB 1817	D39 nanB	Berry <i>et al.</i> (1999)	24.7 ± 4.90	
UB 1912	UB 1830 (pKS80 <i>nanA</i> ⁺)	This study	91.6 ± 4.70	
UB 1913	UB 1817 (pKS80 <i>nanA</i> ⁺)	This study	56.9 ± 5.40	
UB 1915	UB 1816 (pKS80 <i>aad9 nanA</i> ⁺)	This study	44.6 ± 6.30	
UB 1916	UB 1817 (pKS80 <i>aad9 nanA</i> ⁺)	This study	54.0 ± 7.10	
UB 1341	pavA1::cat	Holmes <i>et al.</i> , 2001	46.0 ± 3.23	
UB 1920	UB 1341 (pKS80 nanA ⁺)	This study	76.9 ± 4.50	
UB979	S. pneumoniae 25925 serotype 23F	PHLS (meningitis) CSF	28.0 ± 2.0	
UB1050	Streptococcus pneumoniae 45567 serotype 14	PHLS (pneumonia) blood	158 ± 24.7	
UB 1504	S. gordonii DL1 (wild-type)	D. LeBlanc	<2.0	
UB 1863	UB 1504 (pKS80 <i>nanA</i> ⁺)	This study	61.9 ± 3.71	
UB 1545	S. gordonii hsa::aphA-3	Kerrigan et al., 2007	<2.0	
UB 1897	UB 1545 (pKS80 <i>nanA</i> ⁺)	This study	66.6 ± 7.41	
UB 754	L. lactis MG1363 (wild-type)	M. Gasson	<2.0	
UB 2132	<i>L. lactis</i> (pKS80)	This study	<2.0	
UB 2138	L. lactis (pKS80 nanA ⁺)	This study	41.3 ± 12.0	

 $A_{405} = 1.0$ is equivalent to 0.2 U neuraminidase/10¹⁰ cells.

was unaltered in NanA activity (Table 1) and so the colonization-deficient phenotype of the *pavA* mutant remains unexplained for now (Kadioglu *et al.*, 2010).

nanA complements in *trans* the *nanA* and *nanB* mutant phenotypes

An explanation for the *nanB* mutant phenotype data above was that mutation of *nanB* affected expression

of NanA. To test this possibility, and to also complement the *nanA* mutant, plasmids pKS80 (*nanA*⁺) or pKS80 (*aad9 nanA*⁺) were introduced into the *nanB* or *nanA* mutant strains, respectively. Complementation of the *nanA* mutant produced cells with almost fully restored neuraminidase activity (Table 1; Fig. 4 A), and increased levels of adherence to A549 cells, but only to approximately 50% of wild-type levels (Fig. 4B). Complementation of the *nanB* mutant also



Figure 4 Effect of complementation with *nanA* in trans on (A) enzymic activity, or (B) adherence to A549 cells, of *Streptococcus pneumoniae* D39 wild-type, *nanA* mutant, *nanB* mutant and *S. gordonii* DL1. (–) control pKS80 vector only, (+) corresponding strain carrying plasmid pKS80 *aad9 nanA*+ expressing NanA (see Table 1). Values were obtained from three independent experiments and error bars are \pm SD. **P* ≤ 0.05 relative to respective control.

led to wild-type levels of neuraminidase activity (Fig. 4A) and restored adherence to 58% of wild-type levels (Fig. 4B). This suggested that *nanA* activity *per se* was not the determinant of adherence. Over-expression of *nanA* in the wild-type strain D39 (Fig. 4 A) did not result in increased levels of adherence, suggesting that there may be a threshold of NanA activity above which adherence is limited by other factors.

We also expressed the nanA gene in the oral bacterium S. gordonii DL1, which is normally devoid of neuraminidase activity (Fig. 4A). S. gordonii adheres to sialic acid residues on the surfaces of epithelial cells, and platelets, through the activity of sialic acidbinding adhesin Hsa (Jakubovics et al., 2005a,b; Kerrigan et al., 2007; Jakubovics et al., 2009). It was hypothesized therefore that removal of sialic acid residues by NanA would result in reduced levels of adherence. Indeed, S. gordonii DL1 expressing nanA was \sim 30% reduced in adherence levels to epithelial cells compared with wild-type (Fig. 4B). Moreover, expression of nanA in hsa mutant strain UB1897 had no effect on adhesion levels to epithelial cells (data not shown). These results demonstrated that neuraminidase activity, expressed on the bacterial cell surface, was able to modulate adherence levels by removing sialic acid receptors from the host cell surface.

In situ complementation of nanA mutant

To take advantage of the observation that NanA expressed on the surface of a heterologous cell was efficient at removing sialic acid molecules available for adhesion, we expressed nanA in Lactococcus lactis. This bacterium does not produce neuraminidase activity, and cells do not adhere to epithelial cells or to human tissue proteins (Jakubovics et al., 2009). When A549 cells were pre-incubated with L. lactis, subsequent adherence levels of pneumococcal wildtype or nanA mutant strains were slightly reduced, although not statistically significantly (Fig. 5). However, when A549 cells were pre-incubated with L. lactis expressing NanA activity, adherence levels of the nanA mutant were restored to wild-type (Fig. 5). This suggested that enzymic activity, applied to the epithelial cells in situ prior to adherence assay, was sufficient to expose receptors necessary for adherence of the pneumococcal nanA mutant. However, the effect



Figure 5 Effect of pre-incubating A549 cell monolayers with *L. lactis* MG1363 cells expressing NanA on subsequent adherence of *S. pneumoniae* D39 or *nanA* mutant. Cultured cells were incubated for 30 min at 37 °C with *L. lactis* pKS80 (UB2132) (vector control), *L. lactis* expressing NanA (UB2138), or no *L. lactis* cells. Epithelial cells were then washed and assayed for ability to support adherence of *S. pneumoniae* D39 or *nanA* mutant cells by viable plate count. Values were obtained with triplicate samples from two independent experiments and error bars are ±SD. *P* values are as indicated.

seemed to be specific to NanA, because incubation of epithelial cells with clostridial sialidase did not complement the *nanA* mutant (data not shown). Therefore it is concluded that the *nanA* mutation does not affect expression of adhesins, but affects the ability to specifically expose the necessary receptors for adherence.

NanA expression modulates biofilm formation

To test the role of NanA in biofilm formation, which results from initial adherence, we utilized salivacoated glass cover slips for biofilm growth. The *nanA* and *nanB* mutants were both deficient in biofilm production, measured as biomass, compared with wildtype (Fig. 6A). The growth rates of the *nanA* or *nanB* mutant strains were similar to wild-type D39 in the medium (THY) utilized for biofilm formation. Complementation of the mutants with the corresponding *nanA*-expressing plasmids led to significantly increased biomass in each case, and restored biomass production by the *nanB* mutant to wild-type levels (Fig. 6A). Biofilm formation by *S. pneumoniae*



Figure 6 NanA expression promotes biofilm formation. Panel A, effects of *nanA* or *nanB* mutations, or 10 mM deydro-NeuNAC (dHN) on *Streptococcus pneumoniae* biofilm formation (biomass), and of complementation of mutants with NanA⁺. * $P \le 0.05$ relative to D39, ** $P \le 0.05$ relative to respective non-complemented mutant. Panel B, effects of over-expression of NanA from pKS80 *nanA*⁺ on biofilm formation by *L. lactis* MG1363 (UB2138), in the presence or absence of 10 mM dHN, or by *S. gordonii* DL1 (UB1897). Results were obtained from three independent experiments and error bars are ±SD. * $P \le 0.05$ relative to respective wild-type (wt). Panel C, fluorescence microscopy images of biofilms formed on saliva-coated glass surfaces by (a) *L. lactis* MG1363 (pKS80), (b) *L. lactis* NanA⁺ (UB2138), (c) *S. gordonii* DL1, and (d) *S. gordonii* Nan⁺ (UB1863).

D39, measured as biomass, was 45% reduced in the presence of neuraminidase activity inhibitor 10 mm-NeuNAc (Fig. 6A). This is in keeping with previous observations of the inhibitory effects of neuraminidase enzyme inhibitors on pneumococcal biofilm formation (Parker *et al.*, 2009).

To then test the direct role of *nanA* in biofilm formation, *L. lactis* or *S. gordonii* strains expressing *nanA* were compared with corresponding wild type strains in respect of biofilm biomass and architecture. Expression of *nanA* conferred on *L. lactis* the ability to form a biofilm (Fig. 6B), without which the organism formed only a sparse covering of the substratum (Fig. 6C, panel a). Expression of *nanA* clearly promoted formation of a dense layer of *L. lactis* cells that tended to form clusters (Fig. 6C, panel b). In contrast to the results for *S. pneumoniae* D39, biofilm formation by *L. lactis* expressing NanA was unaf-

fected in the presence of 10 mm dehydro-NeuNAc (Fig. 6B), suggesting that enzymatic activity was not the main reason for promotion of biofilm formation. In addition, biofilm formation by S. gordonii was significantly enhanced by expression of nanA (Fig. 6B, Fig. 6C, panels c and d), with dense clusters of cells appearing to be contained within a matrix. These observations generally are in contrast to those obtained for adherence, where nanA expression did not confer epithelial cell adherence properties upon L. lactis, and resulted in decreased levels of S. gordonii adherence (Fig. 4B). This suggests that under the conditions employed for biofilm formation by NanA-expressing surrogate hosts, NanA acts as an adhesin promoting formation of the microbial community. Overall, these data support the evidence that NanA contains at least two functional domains, one for enzymic activity and another for adherence.

DISCUSSION

This study was undertaken to provide in vitro data to account for, or otherwise, the defects of S. pneumoniae nanA and nanB mutants in their abilities to colonize the murine upper respiratory tract and to translocate to the lungs (Manco et al., 2006). We show here that under our assay conditions the nanA and nanB mutants are deficient in adherence to four epithelial cell types. This would tend to support the hypothesis that one of the reasons that the nanA and nanB mutants were colonization deficient in vivo was because of attenuation in ability to adhere to epithelial cells. Our observations are in contrast to those of King et al. (2004, 2006) who reported that their strain D39AnanA mutant was unaffected in ability to colonize rodent models or adhere to two human epithelial cell lines. The conflicting results with respect to in vivo studies may be related to the use of different inocula sizes and animal backgrounds (Manco et al., 2006). With regard to the discrepancies between our adherence data and those of King et al. (2004), these could be due to our different assay techniques (Gou-Id & Weiser, 2002). In addition to suggesting a role for nanA in adherence, we have shown that nanA is necessary for in vitro microbial community development. Thus, nanA may impact on the ability of pneumococci to form biofilms or aggregates for efficient colonization of the nasopharynx, a prerequisite for translocation to the lungs (Orihuela et al., 2004;

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Oggioni *et al.*, 2006). An analogous effect was recently demonstrated for surface protein PsrP, which appears to induce aggregation of pneumococcal cells and enhance colonization of the lungs (Sanchez *et al.*, 2010). In addition, it seems likely that the inability of the *nanB* mutant to colonize efficiently *in vivo* might be related, at least in part, to reduced expression of NanA activity.

There have been a relatively large number of studies on pneumococcus NanA because of its proposed association with infection, and the ability of a NanA vaccine to protect against pneumococcal challenge (Long et al., 2004). However, the role of NanA in epithelial cell interactions has not been fully understood. The enzyme has been shown to possess a catalytic domain, within the C-terminal region, and an N-terminal adhesion domain that is proposed to be lectin-like in binding endothelial cell surface receptors (Uchiyama et al., 2009). In studies with brain microvascular endothelial cells it was shown that the NanA lectinlike domain promoted chemokine induction and pneumococcal invasion (Banerjee et al., 2010). However, in our experiments, recombinant L. lactis expressing NanA were not enhanced in adherence levels to epithelial cells, suggesting a fundamental difference in functioning of NanA in mediating epithelial vs. endothelial cell adherence. Incubation of Detroit 562 epithelial cells with L. lactis expressing NanA led to modification of the epithelial cell surface such that subsequent adherence levels of the nanA mutant were restored to wild-type. These observations are therefore in keeping with the notion that epithelial cell adherence requires NanA enzymic activity in order to unmask receptors to which pneumococci are able to adhere. In addition, it would appear that epithelial cell adherence by pneumococcus is not driven by the lectin-like domain of NanA, as has been shown for adherence to endothelial cells (Uchiyama et al., 2009).

Other results presented here suggest that NanA functions as an epithelial adherence factor for pneumococcus as a result of enzymatic activity. Thus, over-expression of NanA in *nanA* and *nanB* mutant backgrounds led to increased adherence levels, and adherence to epithelial cells was blocked by pretreatment of pneumococcal cell extracts with enzyme inhibitor Dehydro-NeuNAc. Furthermore, expression of NanA in *S. gordonii*, which produces a sialic acidbinding adhesin Hsa (Kerrigan *et al.*, 2007), led to

30% reduced adherence levels to epithelial cells. Taken together, these observations suggest that sialidase activity may be important for survival of S. pneumoniae within the niche. Streptococcus pneumoniae, unlike S. gordonii, does not express a sialic acid-binding adhesin but, as discussed above, may rely upon decryption of sialic acid glycoconjugates by NanA to expose novel receptors. Concomitantly, the sialidase activity would remove sialic acid receptors utilized by other bacteria for adhesion, such as S. gordonii (Bensing et al., 2004; Takamatsu et al., 2006) and Streptococcus sanguinis (Cowan et al., 1987; Neeser et al., 1995; Plummer et al., 2005), thus potentially helping to relieve competition in colonization of upper respiratory tract surfaces by commensal streptococci.

A summary of the multi-functional and temporal roles of NanA in pneumococcal infection is as follows. Activity of NanA is suggested from our results as necessary for exposure of cryptic receptors for adherence of S. pneumoniae to epithelia, and then subsequently for community (biofilm) formation (Soong et al., 2006). This process could be promoted by continued release of free sialic acid (Trappetti et al., 2009) and up-regulation of NanA production (LeMessurier et al., 2006; Oggioni et al., 2006; Williamson et al., 2008). The role of NanA in biofilm formation has been previously noted (Parker et al., 2009) where nanA mutants were unable to form cellular aggregates. Since inhibition of NanA enzymatic activity does not affect biofilm formation by L. lactis expressing NanA, it is proposed that NanA functions principally as an adhesin under these conditions. The lectin-like domain of NanA may then promote internalization of pneumococci by endothelial cells, systemic spread, induction of pro-inflammatory responses by leukocytes (Chang et al., 2012) and crossing of the blood brain barrier (Uchiyama et al., 2009). Such a multi-faceted protein would clearly be a highly desirable component of an effective vaccine.

An unexplained observation following from this study is that inactivation of *nanB* resulted in approximately 50% reduction in NanA activity. Previous data have shown that inactivation of *nanB* did not affect transcription levels of *nanA*, and *vice versa* (Manco *et al.*, 2006). Therefore, to explain the reduction in NanA activity in *nanB* mutants we hypothesize that there must be some form of control operating at the translational level on NanA expression. Complementation of the *nanA* mutant with plasmid pKS80 *nanA*⁺,

where *nanA* expression is under plasmid-based control sequences, including the ribosome binding site, would naturally circumvent any translational controls operating on *nanA*. This might explain why NanA activity could be restored in the *nanB* mutant complemented *in trans* with pKS80 expressing NanA. It should be noted that the assay utilized for NanA activity would not detect NanB (Xu *et al.*, 2011).

In conclusion, these data provide a clearer understanding of multiple activities of NanA in pneumococcal infections. The experiments were conducted with encapsulated strain D39 cells, suggesting that NanA activity must be projected at the capsule periphery in order that enzymic activity can access the substrate. Moreover, the adhesins that come into play for receptor recognition following desialylation must also be surface exposed. The precise nature of these adhesins is not known and is currently under investigation. These might provide additional targets for vaccines or pharmaceuticals to control epithelial cell colonization by *S. pneumoniae*.

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