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# Exposure of Thomsen-Friedenreich Antigen *in Streptococcus pneumoniae* Infection is Dependent on Pneumococcal Neuraminidase A<sup>\*\*</sup>

**Mamie T. Coats**<sup>a,\*</sup>, **Trudy Murphy**<sup>b</sup>, **James C. Paton**<sup>c</sup>, **Barry Gray**<sup>d</sup>, and **David E. Briles**<sup>a,e</sup> <sup>a</sup> Department of Microbiology, University of Alabama at Birmingham, Birmingham, Alabama

<sup>b</sup> Office of the Director, Division of Bacterial Diseases, National Center for Immunization and Respiratory Diseases, Centers for Disease Control and Prevention, Atlanta, GA

<sup>c</sup> Research Centre for Infectious Diseases, School of Molecular and Biomedical Sciences, University of Adelaide, Adelaide, South Australia

<sup>d</sup> Department of Pediatrics, University of Illinois at Peoria, Peoria Illinois

<sup>e</sup> Department of Pediatrics, University of Alabama at Birmingham, Birmingham, Alabama

### Abstract

Pneumococcal hemolytic uremic syndrome is recognized in a small portion of otherwise healthy children who have or have recently had Streptococcus pneumoniae infections, including severe pneumonia, meningitis, and bacteremia. As in other types of hemolytic uremic syndrome (HUS), pneumococcal HUS is characterized by microangiopathic hemolytic anemia, and thrombocytopenia, usually with extensive kidney damage. Although not demonstrated in vivo, the pathogenesis of pneumococcal HUS has been attributed to the action pneumococcal neuraminidase exposing the usually cryptic Thomsen-Friedenreich antigen (T-antigen) on red blood cells (RBC), and kidney glomeruli. We evaluated the effect of pneumococcal infection on desialylation of RBC and glomeruli during pneumococcal infections in mice. Following intravenous infection with capsular type 19F pneumococci, CFU levels exceeding 1000 CFU /mL blood by the third day were significantly more likely to result in exposed T-antigen on RBC than lower levels of bacteremia. In a pneumonia model, significantly more T-antigen was exposed on RBC in mice treated with penicillin than in those receiving mock treatment. Utilizing mutant pneumococci, we demonstrated that neuraminidase A but not neuraminidase B was necessary for exposure of T-antigen on RBC in vivo. Thus, pneumococcal neuraminidase A is necessary for the exposure of T-antigen in vivo and treatment with penicillin increases this effect. Interestingly, NanA<sup>-</sup> pneumococci were found in the blood in higher numbers and caused more deaths than wild type, NanB<sup>-</sup>, or the NanA<sup>-</sup>/NanB<sup>-</sup> pneumococci.

<sup>\*\*</sup> The findings and conclusions in this report are those of the author(s) and do not necessarily represent the official position of the CDC (Centers for Disease Control and Prevention).

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<sup>\*</sup>Corresponding Author David E. Briles, 1530 3<sup>rd</sup> Ave South, Birmingham AL 35294-2170, (205) 934-6595 (205), 934-9256 (fax), dbriles@uab.edu.

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#### Keywords

*Streptococcus pneumoniae*; Hemolytic uremic syndrome; Thomsen-Friedenreich antigen; Neuraminidase

#### 1. INTRODUCTION

Hemolytic uremic syndrome [1] is characterized by microangiopathic hemolytic anemia, acute renal failure, and thrombocytopenia [2–5]. Pneumococcal HUS is a relatively uncommon complication of invasive infection with *Streptococcus pneumoniae* and is classified as "infection-induced" HUS along with HUS caused by *Escherichia coli* O157:H7 and other bacteria producing shiga-like toxins [6]. The original case series describing HUS by Gasser et al. in 1955 included 2 infants with pneumonia, presumably pneumococcal pneumonia [7]. Pneumococcal HUS is most commonly seen in patients with pneumococcal pneumonia and/or meningitis and unlike HUS caused by *E. coli*, pneumococcal HUS occurs in younger age groups (most patients are less than 2 years), and is more likely to require renal dialysis and prolonged hospitalization. Pneumococcal HUS is estimated to comprise ~5 percent of all HUS cases in children and 0.4% - 0.6% of all invasive pneumococcal infections, but this figure is probably an underestimated because of lack of awareness [3, 4, 8].

A hypothesis for the pathogenesis of pneumococcal HUS was proposed by Klein in 1977, who found Thomsen-Friedenreich antigen (T-antigen) exposed on red blood cells (RBC) and in glomeruli of two children who died of pneumococcal pneumonia and sepsis [9]. The T-antigen is a disaccharide (Gal $\beta$ 1-3GalNAc $\alpha$ 1-Ser/thr) that forms the core structure of *O*-linked mucin type glycans. The T-antigen is "cryptic", normally being hidden by a terminal sialic acid [10] and is found on many cell types including RBC, platelets, glomerular capillary endothelium and renal tubular epithelium.

The neuraminidase from *S. pneumoniae* is hypothesized to expose T-antigen to naturally occurring IgM antibodies by the cleaving terminal sialic acid. Specifically these neuraminidases cleave sialic acid-containing substrates with  $\alpha$ -2,6 and  $\alpha$ -2,3 linkages of N-acetylneuraminic acid to galactose, and  $\alpha$ -2,6 linkage to N-acetyl-galactosamine [11]. The Coomb's test is usually positive in pneumococcal HUS patients, and T-antigen exposure on RBC and other cells can be detected in human and mouse samples using peanut lectin from *Arachis hypogaea* [12, 13]. The finding of neuraminidase activity in *S. pneumoniae* patient isolates is suggestive but not specific for pneumococcal HUS [3]. The neuraminidase activity has been assumed but not proven to be due to pneumococcal neuraminidase [14–17]. The involvement of antibody to T-antigen in pneumococcal HUS pathology is controversial [8]. Although some investigators speculate that a heavy bacterial load with pneumococcal may contribute to the development of HUS, it remains unclear why only a small percentage of invasive pneumococcal infections are complicated by HUS [14].

The *S. pneumoniae* genome encodes 3 neuraminidases: NanA, NanB and NanC [18]. The majority of isolates express NanA and NanB. NanC has high sequence identity to NanB but is not well characterized otherwise. Virtually all strains of *S. pneumoniae* possess the gene for neuraminidase A, and exhibit neuraminidase activity [18–21]. NanA has been shown to be important in the adherence to tracheal epithelium and nasopharyngeal colonization [22, 23] and binding/invasion of the blood-brain barrier [24]. NanA is thought to promote nasal colonization by removing sialic acid from cell surfaces thus exposing receptors to which pneumococci are able to bind [23, 25, 26]. Neuraminidase is also thought to act on mucin,

glycolipids, and glycoproteins [27]. It can remove sialic acid from molecules such as IgA and lactoferrin and has been hypothesized to interfere with protection mediated by these molecules [28]. In most pneumococci, NanA is covalently attached to the peptidoglycan on the surface of *S. pneumoniae* by Sortase A, which recognizes the LPETG cell wall attachment motif of NanA [29–31]. TIGR4 is a commonly used strain in which a naturally-occurring mutation prevents cell wall attachment and the protein is released from the cell surface [31].

Unlike NanA, NanB has no attachment motif and is released into the environment [32]. NanB acts as an intramolecular trans-sialidase specifically cleaving  $\alpha 2$ –3 linked sialic acid substrates [33]. Both NanA and NanB have been reported to play roles in the development of upper and lower respiratory tract infection and sepsis [34]. In animal studies, neuraminidase has been reported to play a role in virulence at the mucosal surface [23] and in systemic disease [34]. Although evidence suggests that neuraminidase is not well expressed by pneumococci during systemic infection, immunization with NanA does offer some protection against sepsis following pneumococcal challenge [28, 35, 36].

We used animal models and in vitro assays to evaluate the ability of NanA and NanB to act in vivo in mice to expose T-antigen. We studied capsular type 19 and 14 strains because they are among the most common capsular types reported in cases of pneumococcal HUS [5]. Our findings demonstrated that during invasive infection in mice enough neuraminidase is made to expose T-antigen on host surfaces of RBC and glomeruli. By using mutant pneumococci we were able to show that NanA, but not NanB, was required for in vivo desialation of host cells in vivo. We also examined the possibility that penicillin, which results in the lytic death of pneumococci, may release enough neuraminidase from the cell wall to enhance desialation of host RBC.

#### 2. RESULTS

# 2.1. Kidney tissue in CBA/N mice is sensitive to the neuraminidase activity of *S. pneumoniae*

In the present study we found that intravenous infection of CBA/N mice with EF3030, HUS05, TJ0893, HUS03 and BG8090 strains of *S. pneumoniae* resulted in the exposure of T-antigen in the kidney along capillaries in the glomeruli, as demonstrated histologically using peroxidase labeled peanut lectin (Figure 1). Glomeruli with exposed T-antigen were located throughout the kidney. Using this qualitative assay differences were not observed in the ability of different strains to expose T-antigen in glomeruli. As expected, infection with NanA<sup>-</sup> mutants of EF3030 did not expose T-antigen on glomeruli (data not shown). The negative control for this study was examination of glomeruli of mice injected with lactated ringers instead of pneumococci. Tantigen was not exposed on the glomeruli of these "mock infected" mice (data not shown).

#### 2.2. Bacteremia is sufficient to cause exposure of T-antigen on mouse RBC

Preliminary studies showed that intravenous infection with ~ $10^7$  CFU of the strains used in the study above generally led to self-resolving bacteremia in CBA/N mice. Peak levels of bacteremia generally occurred by day 3, and by day 5, when the mice were euthanized, bacteremia was  $\leq 100$  CFUs/mL in all mice. Three to five-fold higher numbers of CFU invariably lead to instant sepsis and death within 24 hours. As a result we only monitored T-antigen exposure the infections with  $10^7$  CFU.

The maximum CFUs observed for each mouse during the first 72 hours following infection is shown in figure 2. For 23 of the 26 mice infected with these 5 strains that had bacteremia of  $\geq$ 1000 CFUs/mL exposure of T-antigen was observed; none of the 23 mice with <1000

CFU/mL exhibited T-antigen on their RBC by our assay (P<0.0001; Fisher exact test). Of the **5** strains examined HUS03 appeared to differ from the group as a whole however (P<0.04; Fisher exact test), in that  $\geq$ 200,000 CFU/mL appeared to be its threshold for exposure of T-antigen.

These results raised the possibility that differences in NanA levels produced by the different bacteria were responsible for the difference in the threshold of bacteremia associated with T-antigen exposure between HUS03 and the other strains examined. To examine this possibility growth curves were conducted with the strains in our standard THY broth and levels of neuraminidase activity at various time points. The neuraminidase activity observed for HUS03 in vitro at all phases of growth was indistinguishable from that of the other strains (data not shown). These findings, however, do not rule out the possibility that the strains still might differ in neuraminidase production during an in vivo during an infection.

#### 2.3. Treatment with penicillin increased the chance of T-antigen exposure

Since, neuraminidase A is cell surface attached it was possible that the lysis of pneumococci caused by treatment with antibiotics could release neuraminidase and make it more able to pass from the lung into the blood. To test this possibility mice were infected with EF3030 via nasal aspiration under conditions where the pneumococci were not detected in the blood [37]. After 5 days of antibiotic treatment, the bacterial load in the lungs was almost 100-fold lower in penicillin treated mice than in Ringer's solution treated mice (P=0.0001, Mann-Whitney test) (Figure 3). The bacterial load in the untreated group of mice euthanized at 24 hours was higher than that of Ringer's-treated mice on day 6 (Figure 3) indicating that the EF3030 infection in mice was self resolving as expected [37].

However, the infected mice receiving 6000U of penicillin were statistically more likely to have T-antigen exposed on their RBC (P=0.0131, Fisher's exact test) than infected mice receiving only Ringer's solution. Of the 39 infected mice only 3 were found to be bacteremic (CFUs > 10<sup>1.49</sup> CFU/mL). These three mice were excluded from statistical analysis of T-antigen exposure. Our results suggest that the penicillin induced autolysis of pneumococci [38, 39] resulted in neuraminidase release from the bacteria in the lungs, which reached the circulation where it was able expose T-antigen exposure on RBC.

#### 2.4. Neuraminidase deficient strains of S. pneumoniae

Parent and neuraminidase deficient strains were grown in THY to O.D. 0.5 at 600 nm. Mutations were verified using PCR, Western blotting and neuraminidase activity measurements (Data not shown).

#### 2.5. NanA exposes T-antigen on RBC in vivo

Intravenous infection with EF3030 resulted in the exposure of T-antigen on circulating RBC (solid symbols, Figure 4). However, infection with strains deficient in NanA alone (Sam11 and Sam1) or deficient in both NanA and NanB (Sam30 and Sam3) did not result in exposure of T-antigen on circulating RBC (open symbols, Figure 4) following intravenous infection (P=0.002, chi-square analysis). The two strains with mutations in *nanB* but not *nanA* (Sam 20 and Sam 2) exposed T-antigen as efficiently as the wild type strain EF3030. Thus, the expression of NanA but not NanB was required for T-antigen exposure in vivo.

#### 2.6. Effects of NanA and NanB on bacteremia and mortality

Intravenous infections with the *nanA* and *nanB* mutant strains resulted in somewhat higher levels of bacteria in the blood for the mutant strains than for the wild type EF3030. This difference was only statistically significant, for the comparison of the NanA deficient strain with the EF3030 parent (Figure 4). It was also observed that 7 of 10 mice infected with

NanA<sup>-</sup> pneumococci died following lung infection whereas only 0 to 1 of 10 mice infected with either wild type, NanB<sup>-</sup> or NanA /NanB pneumococci died (Figure 4). An earlier study by Manco et al. using the much more virulent D39 strain background observed that both mutations in both *nanA* and *nanB* decreased blood CFUs [34]. The difference in results between their study and ours may be caused by the much greater virulence of D39 versus EF3030 pneumococci, which may affect the relative role of NanA on virulence of the different strains.

#### 3. DISCUSSION AND CONCLUSION

HUS is an unpredictable consequence of *S. pneumoniae* infection of infants and young children. While the incidence of diarrhea associated with Gram-negative HUS is higher than with pneumococcal HUS, patients with pneumococcal HUS require a longer average hospital stay, have a higher mortality, and more frequently require renal dialysis [2, 8]. It has long been assumed that the neuraminidase detected in the blood of patients with pneumococcal HUS is produced by the pneumococci and is responsible for exposing T-antigen in the kidney and on RBC of HUS patients [9]. Our findings provide experimental support for this earlier prediction and demonstrate that NanA, but not NanB is necessary for T-antigen exposure.

Gene activation studies indicate that *nanA* is more strongly expressed in the transparent phase of pneumococci that colonizes the surface of the nasal mucosa than in opaque phase pneumococci responsible for invasive disease [28]. However, NanA is sufficiently expressed during invasive disease that immunity to NanA offers some protection in mice against intraperitoneal-sepsis and intranasal-sepsis following inoculation of *S. pneumoniae* [34, 36].

This study demonstrated that infection of mice with wild-type pneumococci, but not NanAdeficient pneumococci resulted in exposure of T-antigen in vivo. Since the expression of NanA increased the numbers of pneumococci in circulation during the first 72 hours following infection, it was clear that exposure of T-antigen required NanA and not pneumococcal infection *per se*. This conclusion is further supported by the fact that in the study depicted in figure 4 the mice with the highest levels of CFU in the blood were those infected with the NanA pneumococci, and none of them exhibited any exposure of T-antigen on their RBC. NanB, however, had no effect on the exposure of T-antigen in vivo. The activity of NanA but not NanB in the exposure of T-antigen on RBC in vivo was consistent with NanB's reported low enzymatic activity and the fact that the pH optima of NanB (pH 4.5) is well below the pH of blood [32].

The exposure of T-antigen by infecting pneumococci was a general effect of pneumococcal infection, since it was observed with all 5 wild-type strains examined. Several capsular polysaccharide serotypes have been reported in cases of pneumococcal HUS, and case reports suggest many of the patients had a heavy bacterial load, e.g., empyema or more than one site of infection (e.g., pneumonia and meningitis) [4, 8]. We observed that levels of bacteremia exceeding 10<sup>3</sup>/mL in mice were associated with the in vivo exposure of T-antigen on RBC. For one of our strains, however, more CFUs/mL of blood was required to expose T-antigen than for the other four strains. Since we observed no less NanA produced in vitro by this strain, this result suggests that the threshold of bacteremia required for T-antigen exposure might be affected by differences in the activity or amount of NanA produced during infections.

Interestingly, the absence of NanA from EF3030 resulted in higher blood levels of CFU following lung infection, and lead to much higher death rate (Figure 4). While this may seem surprising for an antigen thought of as a virulence antigen, it must be remembered that

selection for invasive disease is probably not necessarily important in the evolution of pneumococci, which are through to be spread largely spread and acquisition of colonization [40]. The greater virulence of the NanA<sup>-</sup> than wild-type EF3030 may be because the NanA<sup>-</sup> bacteria are less adhesive to the host gangliosides and thus more able to spread through tissues [23, 26]. The fact that NanA pneumococci were significantly more virulent than mice lacking both NanA or NanB (Figure 4) is consistent with the claim by Manco et al that NanB has an effect on invasive pneumococcal disease [34].

Our observation that penicillin treatment during lung infection of mice enhanced T-antigen exposure on circulating RBC suggested that neuraminidase released by the autolysis of the pneumococci following antibiotic treatment could reach the circulatory system. Withholding antibiotics during patient treatment is not a viable option in pneumococcal infection. But, one possible approach to minimize T-antigen exposure and resultant risk of HUS might be to give passive antibody to NanA or a neuraminidase inhibitor to infants suffering from sever pneumococcal infections.

A difference in T-antigen exposure, by itself, is probably not the only predictor in the development of HUS [3]. Antibody to neuraminidase develops in most individuals by 2 years of age [41] and antibody titers to T-antigen can be detected in most infants by 6 months of age [8]. Thus, major additional factors predisposing infants to HUS may be variable levels of sensitizing maternal antibodies to T-antigen and the presence of neutralizing maternal antibodies to pneumococcal neuraminidase.

#### 4. METHODS

#### 4.1. S. pneumoniae

Wild-type pneumococcal strains were all clinical isolates (Table 1). Strains EF3030, BG8090, and TJ0893 were capsular types 19F, 19, and 14 [37, 42, 43]. Strains HUS03 (19A) and HUS05 (19F) were collected from cases of pneumococcal HUS [44] in Dallas, Texas, 1997–1998 with IRB approval. Pneumococci were grown to exponential phase in Todd Hewitt broth containing 0.5% yeast extract [45], concentrated by centrifugation when necessary, and were aliquoted and stored in THY containing 10% glycerol at  $-80^{\circ}$ C until use. Strains carrying antibiotic markers were plated on selective blood agar containing appropriate antibiotics.

A NanA deficient strain in the D39 background was made by insertion-duplication mutagenesis and has been described previously [46]. Chromosomal DNA was isolated from this strain and transformed into strain EF3030 to produce a *nanA* mutant of EF3030. Transformations were carried out as described previously [47] and genes were backcrossed into recipient strains three times to eliminate co-transformation with non-contiguous DNA (strains Sam1, Sam11). Transformants were selected based on erythromycin resistance and verified with PCR. EF3030 NanB deficient strains were also constructed using the insertion-duplication mutagenesis technique (Sam2, Sam20). EF3030 derivatives lacking NanA and NanB were constructed by transforming EF3030 with DNA from an existing *nanA/nanB* double mutant into EF3030 to produce two independent strains containing both the NanA and NanB mutations (Sam3, Sam30) [34]. The double transformants were selected using erythromycin alone; as *nanA* and *nanB* are closely linked on the chromosome [29] and co-transformants containing both mutants could easily be recovered. The presence of both mutations in the new strains was confirmed by PCR analysis.

#### 4.2. Mice

Female CBA/CaHN-*Btkxid*/J (CBA/N) mice were purchased for Jackson Laboratory, Bar Harbor, Maine. Mice were 8–12 weeks old at the time of the experiments. Animal studies were done under approval from University of Alabama at Birmingham IACUC 041206026.

#### 4.3. Intravenous infection of CBA/N mice with S. pneumoniae

Bacteremia/sepsis was induced by intravenous infection  $\sim 10^7$  CFUs of the indicated strains as described [48]. Eight to 11 mice were used in each group. Blood (0.07 ml diluted immediately with 500µl of 1% BSA in PBS) was collected every 24 hours by retro-orbital puncture under isoflurane (Minrad Inc. Buffalo, NY) for the first 3 days post infection, and when indicated on day 5. Serial dilutions of the blood were plated on blood agar plates and CFUs enumerated [48]. Mice were euthanized 5 or 7-days post infection as indicated and kidneys were harvested where indicated.

#### 4.4. Detection of T-antigen exposure on RBC

To detect T-antigen exposure on RBC, diluted blood was collected as described and the RBC pelleted. The pelleted RBC were suspended to 2% in physiologic saline. One volume of RBC was incubated with 2 volumes of *Arachis hypogea* lectin, peanut lectin, (Gamma Biologicals, Houston, TX) for 5 minutes at room temperature. Peanut lectin binds Galβ-1-3GalNAc. The mixture was centrifuged to pellet the RBC. RBC were resuspended and read macroscopically for agglutination. Detectable RBC agglutination following centrifugation was indicative of T-antigen exposure (T-antigen<sup>+</sup>). RBC that failed to show detectable agglutination were considered negative for T-antigen exposure (T-antigen<sup>-</sup>). Negative agglutination was verified using a 1.5X magnifying glass.

#### 4.5. T-antigen exposure in the kidney

For histology, organs from infected mice were fixed in 10% formalin, dehydrated, and embedded in paraffin. Four-micron thick sections were cut. Sections were deparaffinized in 2 changes of xylene and rehydrated through graded ethanol solutions ending in deionized water. Exposed T-antigen was detected by incubating the sections with 1mg/mL horse radish peroxidase (HRP)-labeled peanut lectin (Sigma, St. Louis, Missouri) for 1 hour at room temperature. Following incubation slides were incubated with 3-3' diaminobenzidine tetrahydrochloride (DAB) (Sigma, St. Louis, Missouri) at room temperature for 30 minutes. Slides were dehydrated using ethanol, mounted and observed.

#### 4.6. The effect of penicillin on the exposure of T-antigen by infecting pneumococci

A mouse model of pneumococcal pneumonia was used [47]. Mice were anesthetized to promote aspiration prior to being inoculated intranasally with  $10^7$  CFUs of EF3030 in 40 µl Ringer's Injection solution (Abbott labs, Chicago). Beginning 24 hours after infection, mice were treated intraperitoneally with 6000U penicillin G or Ringer's solution every 24 hours and were euthanized 6 days post inoculation. A third group of mice received neither penicillin nor Ringer's solution and were euthanized 24 hours after infection. The purpose of the third group was to quantitate the levels of bacteria in the lungs prior to any treatment. All groups of mice were monitored for survival every 8 hours until euthanized. When each mouse was euthanized lungs, nasal wash, and blood were collected. Organs and blood were plated on blood agar plates containing gentamicin (4µg/mL).

#### 4.7. Neuraminidase activity

Neuraminidase activity was detected on logarithmic phase cultures in a quantitative assay using the substrate, 2'-(4-methylumbelliferyl)-a-D-N-acetylneuraminic acid (MUAN) (Sigma Chemical) as described by Lock et al [20]. The substrate, 2'-(4-methylumbelliferyl)-

a-D-N-acetylneuraminic acid (MUAN) (Sigma Chemical), is prepared as a 0.35% solution in water. Bacteria were grown to an optical density of 0.5 at 600nm. Cultures contained approximately  $1 \times 10^6$  CFUs/mL bacteria. Cells were centrifuged pelleted. Cell pellets and culture supernatants were collected. Cell pellets were lysed using a solution containing 0.1% sodium deoxycholate, 0.01% sodium dodecyl sulfate, and 0.15M sodium citrate) at stored at

sodium deoxycholate, 0.01% sodium dodecyl sulfate, and 0.15M sodium citrate) at stored at -20°C until time of assay. For neuraminidase activity measurements, equal volumes of substrate and cell lysate or supernatant were mixed. Mixtures were incubated for 5 minutes at 37°C at pH 7.0. The reaction was stopped by the addition of 10 volumes of 0.5 M sodium carbonate buffer (pH 10.5). The amount of 4-methyl-umbelliferone (MU) released was measured (excitation wavelength of 355 nm and emission wavelength of 460 nM) [49]. Measurements were made and quantitated by comparison to the optical density of a known concentration of MU.

#### 4.8. Statistical analysis

For statistical analysis, Mann Whitney, chi square analysis, Kruskal-Wallis, and Duns multiple comparison were used as indicated.

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#### Figure 1.

Exposure of T-antigen on glomeruli during infection with *S. pneumoniae*. Mice were infected intravenously with *S. pneumoniae*  $1 \times 10^7$  CFU of EF3030. Fixed kidneys were reacted with HRP-labeled peanut lectin. Control kidneys from mice injected with ringers but no bacteria did not expose T-antigen and no-exposure was seen in kidneys from mice infected with NanA<sup>-</sup> EF3030 (data not shown).

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#### Figure 2.

Bacteremia is sufficient to cause T-antigen exposure during infection. Mice were infected intravenously with ~10<sup>7</sup> CFUs of *S. pneumoniae*. Symbols denote the highest number of CFUs obtained for an individual animal at their 24, 48, and 72 h bleeds following infection. Closed and open symbols indicate animals that become positive or remained negative, respectively, for T-antigen exposure on RBC during five-day infection period. Animals with no detectable bacteremia were assigned a value of  $10^{1.48}$ , the limit of detection of CFUs in the blood. Bacteremia was self-resolving and no mice died following infection. For each strain of bacteria we used the Fisher exact test to look for an association between CFUs greater than  $10^3$  CFUs/mL and T-antigen exposure. The association was significant for all strains at P < 0.02 except for HUS03. Using a multi-row Chi-square contingency table the ability of mice with >10<sup>3</sup> CFUs to exhibit T-antigen on RBC was found to vary among the strains at P < 0.02.



#### Figure 3.

Clearance of *S. pneumoniae* following penicillin treatment increases T-antigen exposure. Mice received lung infections with 10<sup>7</sup> CFUs EF3030 given IN. At 24 hours post infection, animals received penicillin or lactated Ringers treatment every 24 hours until euthanized on day 6, at which time the numbers of CFUs in their lungs were determined. \*Effect of penicillin on exposure of T-antigen in mice, P=0.0131 versus Ringer's solution control by the Fisher's exact test. \*\*Differences in the numbers of CFUs in the lungs of penicillin-treatment versus Ringer's solution control was significant at P≤0.0001, by the Mann-Whitney test. Coats et al.



NS

#### Figure 4.

< 0.01\*

CFU

NanA was required for T-antigen exposure. Groups of 10 mice were infected intravenously with EF3030 or neuraminidase deficient derivatives of EF3030. The mice were bled every 24 hours, until death. Each symbol represents the highest number of CFU observed for each mouse in the first 72 hours. Closed and open symbols indicate animals that were positive or negative for T-antigen on their RBCs during the 5-day period of observation. For each NanA/NanB genotype two independent mutants were examined and the pooled data are shown. Overall the numbers of T antigen positive and negative mice of the different genotypes were found to differ among the groups using a  $2 \times 4$  chi-square analysis (P <0.0001). The differences in CFUs in the blood for the 4 genotypes was significant at P=0.012 by Kruskal-Wallis. With the Duns multiple comparison test only the difference between NanA<sup>-</sup>/NanB<sup>+</sup> and EF3030 was statistically significant (P < 0.05). Mice that died are marked with  $\Box$ . All deaths occurred only on days 2, 3, or 4. The median day of death for mice infected with NanA<sup>-</sup> EF3030 was 3. The single deaths in groups infected with NanA<sup>-</sup>/ NanB<sup>-</sup> and EF303 were on days 4 and 3 respectively. By a 2×4 Chi-Square analysis the distributions of deaths in the four groups was significant by P<0.0005. The NanA<sup>-</sup>/NanB<sup>+</sup> group differed from each of the other groups by Fisher exact test at P < 0.02.

NS

EF3030

#### Table 1

### Strains included in this study

Strain	Capsule type	Phenotype	Derivation	Reference
EF3030	19F	NanA <sup>+</sup> , NanB <sup>+</sup>	Clinical isolate	[37] Otitis Media ~1980
HUS05	19F	NanA <sup>+</sup> , NanB <sup>+</sup>	Clinical isolate	[44] HUS isolate, 1995
BG8090	19	NanA <sup>+</sup> , NanB <sup>+</sup>	Clinical isolate	Meningitis isolate, 1986
HUS03	19A	NanA <sup>+</sup> , NanB <sup>+</sup>	Clinical isolate	[44] HUS isolate, 1995
TJ0893	14	NanA <sup>+</sup> , NanB <sup>+</sup>	Clinical isolate	Pneumonia isolate (AIDS patient), 1993
Sam1	19	NanA <sup>-</sup> , NanB <sup>+</sup>	EF3030	This study
Sam11	19	NanA <sup>-</sup> , NanB <sup>+</sup>	EF3030	This study
Sam2	19	NanA <sup>+</sup> , NanB <sup>-</sup>	EF3030	This study
Sam20	19	NanA+, NanB-	EF3030	This study
Sam3	19	NanA <sup>-</sup> , NanB <sup>-</sup>	EF3030	This study
Sam30	19	NanA <sup>-</sup> , NanB <sup>-</sup>	EF3030	This study