Development of a multicomponent Staphylococcus aureus vaccine designed to counter multiple bacterial virulence factors

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Abbreviations: PMN, polymorphonuclear neutrophil; HA, hospital-acquired; CA, community-acquired; MRSA, methicillinresistance *Staphylococcus aureus*; ClfA, clumping factor A; Fg, fibrinogen; MSCRAMM, microbial surface component recognizing adhesive matrix molecule; MntC, manganese transporter C; IdsB, iron-responsive surface determinant B; CP, capsular polysaccharide; OPA, opsonophagocytosis; OPA, OP assay; FBI, fibrinogen binding inhibition

Staphylococcus aureus is a major cause of healthcare-associated infections and is responsible for a substantial burden of disease in hospitalized patients. Despite increasingly rigorous infection control guidelines, the prevalence and corresponding negative impact of S. aureus infections remain considerable. Difficulties in controlling S. aureus infections as well as the associated treatment costs are exacerbated by increasing rates of resistance to available antibiotics. Despite ongoing efforts over the past 20 years, no licensed S. aureus vaccine is currently available. However, learnings from past clinical failures of vaccine candidates and a better understanding of the immunopathology of S. aureus colonization and infection have aided in the design of new vaccine candidates based on multiple important bacterial pathogenesis mechanisms. This review outlines important considerations in designing a vaccine for the prevention of S. aureus disease in healthcare settings.

The Complexity of Staphylococcal Disease

Staphylococcus aureus (S. aureus) are commensal, Gram-positive bacteria which colonize the nares, axilla, pharynx and other mucosal and skin surfaces of ~30% of human subjects. S. aureus is estimated to be responsible for 20–25% of all healthcare-associated infections,^{1,2} resulting in three times the length of hospital stay and a 5-fold higher risk of in-hospital death for infected patients compared with patients without such infections.³ Furthermore, S. aureus infections can be associated with in-hospital mortality rates of up to 25%.⁴ Although S. aureus is the etiological agent of a diverse number of diseases, including necrotizing pneumonia, septic arthritis and osteomyelitis, 90% of all infections are a result of skin and soft tissue structure breaches.⁵⁻⁷ Historically, S. aureus has been associated mainly with nosocomial infections. Over

*Correspondence to: Kathrin U. Jansen; Email: kathrin.jansen@pfizer.com Submitted: 05/15/12; Revised: 08/09/12; Accepted: 08/17/12 http://dx.doi.org/10.4161/hv.21872 recent decades, however, there has been a dramatic increase in *S. aureus* infection associated with antibiotic resistances throughout the community, most notably in the US.⁸⁻¹¹

Customized Pathogenicity

S. aureus utilizes distinct mechanisms tailored for survival in different microenvironments encountered during host colonization¹²⁻¹⁵ or invasion, such as inhibition of phagocytic uptake and killing,¹⁶⁻¹⁸ dissemination in the bloodstream, and formation of abscesses^{19,20} or biofilms.^{21,22} This versatility arises from the large number of virulence factors which S. aureus deploys in distinct spatial and temporal patterns to optimize its chances of survival.²⁰ These virulence factors include surface proteins that allow adhesion to host components such as fibrinogen (Clumping Factors, ClfAand B) or fibronectin (fibronectin binding proteins, FnBP A and B),^{23,24} and proteins which scavenge nutrients that are normally sequestered in vivo (such as iron-responsive surface determinants, IsdA and B).²⁵ In addition, the bacteria can express an impressive array of factors specifically designed to avoid the immune system, including an anti-opsonic extracellular capsule that protects the bacteria from non-anticapsular polysaccharide antibodies and innate immune components, protein inhibitors of neutrophil chemotaxis and the complement cascade, immunoglobulin binding proteins (such as staphylococcal protein A or Spa) and enzymes which aid its survival within the phagosome of neutrophils (such as the superoxide dismutases SodA and M).²⁶ Most strains of S. aureus also elaborate a number of different invasins (such as hyaluronidase and staphylokinase) and/or toxins (such as enterotoxins A and B, α toxin and Panton-Valentine leukocidin) which promote tissue damage and play an important role in septic shock.27,28

A significant amount of research has explored how *S. aureus* is capable of transitioning from a harmless commensal organism to a life threatening infectious agent.^{15,29-31} It has been observed that people colonized with *S. aureus* are at higher risk of infection

than non-carriers^{32,33} and that those infections usually arise from the colonizing strain,³⁴ yet colonized individuals have a greater chance of recovering from these infections. Interestingly, recovery from a S. aureus infection does not appear to confer immunity to subsequent infections.35 Although somewhat paradoxical, these observations could be interpreted to mean that humans who are naturally exposed to S. aureus through asymptomatic carriage or previous infection may mount a sufficient immune response to the carriage strain to reduce the severity of infection but not to other strains circulating in the hospital or the community. In most cases, S. aureus infection occurs after breaches in the skin or mucosal barriers through wounds, trauma or surgical intervention which give the organism direct access to tissues or the bloodstream.³⁶ Once the skin or mucosa has been penetrated, infection can spread to the blood, causing bacteremia, or disseminate to other sites throughout the body.³⁷ Foreign devices surgically inserted into the body, such as joint prostheses, ventilators or catheters, can also become sites of S. aureus infection; ~20% of infections of implanted devices have been found to be caused by this pathogen.³⁸⁻⁴⁰ The surface characteristics of the device may facilitate bacterial adhesion, thereby increasing the risk of infection. As mentioned above, community-associated infections are increasing in incidence and initial superficial infections as a result of small skin abrasions or other minor skin lesions have the propensity to develop and spread.41

Attack/Counterattack: Mechanisms of Immunity that Control *S. aureus*

Humans can be permanently colonized with S. aureus; as such, there can be a constant interplay between the bacteria and the host immune system. This is supported by the observation that all humans have pre-existing antibodies to S. aureus antigens. However, as discussed below, functional antibodies that facilitate the clearing of staphylococci from the site of an infection, which is through uptake and killing by professional phagocytes, especially neutrophils,^{17,42} or functional antibodies that neutralize virulence factors, are absent in the majority of the human population. The importance of immune clearance is underscored by the increased rates of infection that are observed for subjects with immunological disorders as summarized below; the close link between defects in clearance and the risk of disease provides confidence that if a vaccine generates adequate levels of functional antibodies in subjects with competent effector cells, those individuals are likely to be protected at times when they are at risk of infection.

Polymorphonuclear neutrophils, or PMNs, are the primary cellular defense against staphylococcal infections. These effector cells are highly efficient at killing phagocytosed pathogens by engaging in a complex cascade of cellular events involving multiple defense mechanisms to eradicate bacteria. Phagocytosis is the binding and ingestion of bacteria which can be facilitated by opsonization of the microbial surfaces with antibody and/or complement.⁴³ Neutrophils recognize many molecules produced by *S. aureus* such as lipoteichoic acid (LTA) and CpG DNA which interact with TLRs on the neutrophils to promote phagocytosis.

Additionally, uptake of bacteria is enhanced if they are opsonized by complement and/or antibodies present in serum. Neutrophils express on their cell surfaces both complement receptors that bind complement-opsonized bacteria and Fc receptors that bind the Fc region of antibody-coated bacteria. Once it is bound to the neutrophils, a phagosome is formed to engulf the bacteria. This is followed by neutrophil-initiated mechanisms to kill the bacteria. Phagocytosis activates membrane bound NADPH-dependent oxidase to initiate a "respiratory burst," which generates high levels of reactive oxygen species (ROS), i.e., superoxide radicals, hydrogen peroxide and hydroxyl radicals, to kill microorganisms. In addition, degranulation, the fusion of cytoplasmic granules with the bacteria-containing phagosome, results in the release of anti-microbial peptides, proteases and α -defensins with potent microbicidal activity.²⁶

Individuals with PMN defects have a higher incidence of S. aureus disease. Examples of neutrophil dysfunction include frank neutropenia,44 Chronic Granulomatous Disease, in which neutrophils are unable to generate a functional respiratory burst⁴⁵⁻⁴⁷ and Chediak-Higashi Syndrome, in which individuals have neutrophils with reduced chemotaxis and phagolysosome function.⁴⁸ Subjects with defects in non-antibody-mediated clearance mechanisms, such as those with mutations in the pattern recognition receptor TLR2,49 or defects in the complement pathway⁵⁰ cannot effectively clear staphylococci and thus are more susceptible to S. aureus infections. Individuals with defective STAT3 signaling proteins have also been shown to be more prone to S. aureus infections due to their inability to generate IL-17-producing Th17 cells, which results in diminished neutrophil recruitment and function.⁵¹⁻⁵³ Th17 cells have been shown to be important in response to natural infection with S. aureus, and some studies have shown Th17 responses may contribute to effective vaccine responses.54,55 In addition, S. aureus infection in the absence of vaccination has been shown to induce a Th17/Th1 response.^{56,57} Th17 cells may therefore play an important role in the control of S. aureus infections by serving as an link between neutrophil function and cell-mediated immunity.

Antibody-mediated clearance mechanisms driven by the adaptive immune system, in which antibody-secreting B cells and cytokine-secreting and cytolytic T cells play a key role, are also important in the prevention of staphylococcal disease. For example, individuals with immune defects impairing the ability to produce functional antibodies, such as AIDS,⁵⁸ immature immune systems due to premature birth,⁵⁹ or defects in immunoglobulin production all have increased susceptibility to staphylococcal infections.^{60,61} In males with X-linked agammaglobulinemia, recurrent pyogenic infections, often caused by *S. aureus*, begin to occur within the first year of life, after maternal IgG has been exhausted.⁶² Overall, these findings support the requirement for functional antibodies and a functional neutrophil effector cell function to prevent staphylococcal infections and to prevent disease.

Learnings from Previous Clinical Experience

Several reviews have recently been published which summarize the clinical experience to date for *S. aureus* vaccines and discuss future prospects for the field.^{35,63-67} The following sections describe key learnings from previous clinical experiences that could explain why clinical success may have been elusive to date and how Pfizer has incorporated the acquired knowledge into the design of its current vaccine candidates for the prevention of *S. aureus* infection and disease.

Single vs. multiple targets. Nearly all S. aureus vaccine candidates that have undergone clinical testing to date have targeted single antigens.^{35,65,67} Although this has been a very effective strategy for vaccines against other human bacterial pathogens such as Streptococcus pneumoniae,68 Hemophilus influenzae69 and Neisseria meningitidis,⁷⁰ it has not been effective for S. aureus. NABI conducted clinical studies for the prevention of S. aureus infection in hemodialysis patients using StaphVAX, a bivalent polysaccharide conjugate vaccine using E. coli enterotoxin as a carrier protein.⁷¹ Despite promising preclinical data and demonstrated immunogenicity and safety in humans, administration of StaphVAX did not result in a decrease in the incidence of S. aureus infection in a pivotal phase III trial.⁶⁶ Another more recent example of a vaccine failure using a single S. aureus antigen is Merck's monovalent vaccine based on IsdB, a conserved iron-scavenging protein expressed on the surface of S. aureus⁷² which was identified using a novel screen for antigens which are highly expressed during infection.73 While IsdB showed efficacy in several preclinical murine models of S. aureus infection^{72,74,75} it did not demonstrate efficacy in preventing S. aureus infections in cardiothoracic surgery populations.⁷⁶ These and other clinical failures summarized elsewhere^{64,65,67} underscore the complex nature of S. aureus pathogenicity and suggest that a multiantigen vaccine designed to address major bacterial pathogenesis mechanisms will be required for efficacy. In support of this strategy, a recent preclinical study showed that vaccination of mice with a combination of four S. aureus cell wall-anchored adhesins resulted in 100% survival in a sepsis model of infection whereas immunization with each individual component only resulted in 50-70% protection.77 Accordingly, as described below, Pfizer's vaccine candidate contains multiple components that are designed to combat key pathogenic mechanisms deployed by S. aureus.

Selection of the antigen composition. Capsular polysaccharides are anti-phagocytic and key for immune evasion. Expression of capsular polysaccharides is a common mechanism by which pathogenic bacteria, including S. aureus, evade opsonophagocytosis.³⁶ All S. aureus clinical isolates have the genetic pathway for either capsular polysaccharide (CP) type 5 or 878 and serotyping has shown that the vast majority of S. aureus strains express one or the other.79 Although CP5 and CP8 are comprised of identical monosaccharides (L-FucNAc, D-FucNAc and D-ManNAc), they are serologically distinct due to differences in the linkages between sugars and the sites of O-acetylation.⁸⁰ Encapsulated S. aureus strains are more virulent in bacteremia models compared with capsule-defective isogenic mutants.³ The increased virulence of capsule-expressing strains correlates with increased resistance to in vitro opsonophagocytic (OP) killing by human PMNs.^{81,80} In addition, as shown in Figure 1, S. aureus capsular polysaccharides are expressed early after infection of test animals, indicating

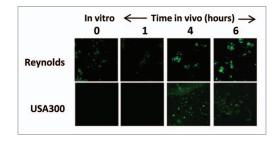


Figure 1. Expression of capsule during infection. Immunofluorescence analysis was performed on the laboratory *S. aureus* strain Reynolds and the USA300 strain CDC3 to determine the timing and degree of capsule expression in a murine infection model of bacteremia. CD1 mice (n = 3) were infected by intraperitoneal injection of approximately 2×10^8 colony-forming units. Immunofluorescence was performed on bacteria harvested at various time points using primary rabbit antibodies and an ALEXA488-conjugated goat- α -rabbit antibody. Both strains express high levels of capsule after four hours of infection (reproduced with permission from (107). Note this is in contrast to several reports that suggest USA300 strains are acapsular due to the lack of observed expression in vitro.^{119,120}

that these structures appear to convey an in vivo survival benefit to the bacteria.

Though capsular polysaccharides provide a good immune evasion strategy for the bacteria, anti- polysaccharide antibodies efficiently bind to bacterial cell surfaces and facilitate phagocytosis. To determine whether S. aureus capsular polysaccharides can induce opsonophagocytic antibody responses, Pfizer conducted preclinical studies with the CP5 and CP8 polysaccharides conjugated to CRM₁₉₇, the same protein carrier also used in other polysaccharide conjugate vaccines such as Prevnar and Prevnar 13®.68 Efficacy of CP5/CP8 conjugates have previously been observed after immunization with the conjugates in preclinical animal models such as murine pylonephritis, rat and rabbit endocarditis and infant rat studies.⁸²⁻⁸⁴ Pfizer recently demonstrated that vaccination with both CP5 and CP8 conjugates in preclinical animal models and in humans resulted in high levels of antibodies that killed S. aureus clinical isolates in opsonophagocytosis assays (OPAs) as described below.

ClfA is a multifunctional virulence factor. Another component that has been selected by Pfizer for inclusion in its multivalent S. aureus vaccine is Clumping Factor A (ClfA). ClfA is a surface adhesin that binds to the C-terminus of the plasma fibrinogen γ chain.^{85,86} This interaction is central to the multifunctional activities of this well-characterized virulence factor. It promotes fibrin cross-linking and mediates the binding of the pathogens to platelets,^{87,88} resulting in thrombus (blood clot) formation. ClfA has also been shown to play a key role in the agglutination of staphylococci in the blood during infection, which leads to thromboembolic lesions in heart tissue and sepsis.⁸⁹ The binding of ClfA to fibrinogen may also subvert opsonophagocytic killing by enhancing the binding of complement factor I to ClfA on the bacterial surface, which in turn promotes cleavage of opsonin C3b into inactive fragments.^{90,91} The fibrinogen binding activity of ClfA is linked to the ability of *S. aureus* to cause disease, as *S.* aureus strains with ClfA point mutations that prevent fibrinogen binding show reduced virulence.⁹² The ClfA antigen is detected in over 99% of *S. aureus* clinical isolates and is highly conserved (> 88% sequence identity).

The importance of ClfA as a virulence factor is illustrated by its ability to confer virulence, independent of other surface antigens, to an otherwise harmless strain of bacteria, Lactococcus lactis. L. lactis is able to process recombinant S. aureus ClfA, expressing it on its own surface at levels similar to that seen on S. aureus.93 Heterologous expression of the ClfA adhesin in L. lactis allows the bacterium to then successfully colonize damaged valves in an experimental rat endocarditis model.⁹⁴ Recent unpublished data from Magnus Hook's lab (U. Texas A and M) demonstrate that a L. lactis ClfA-expressing strain is capable of killing mice in a mouse bacteremia survival model (data not shown). In contrast, control strains expressing a ClfA mutant that cannot bind to fibrinogen (ClfAm) or that harbor the empty plasmid vector are completely avirulent (p = < 0.001 compared with L. lactis with intact ClfA). Therefore, the ability of S. aureus cells to bind to fibrinogen via ClfA is an important virulence mechanism of the bacteria. This phenotype has been captured in the Fibrinogen Binding Inhibition (FBI) Assay, which, as described below, measures the ability of antibodies to prevent S. aureus from binding to human fibrinogen.

Numerous studies suggest that ClfA is a promising component of an efficacious S. aureus vaccine. Preclinically, it was shown to be protective in murine models of arthritis, sepsis and endocarditis.^{92,95,96} A particularly relevant study recently demonstrated the benefit of targeting both CP5 and ClfA antigens in a mouse mastitis model, in which passive immunization with antibodies to the two antigens had an additive effect on reducing bacterial burden.97 In the clinic, ClfA was indirectly evaluated in studies with Veronate, a pool of plasma-derived, donor-selected, polyclonal antistaphylococcal human IgG with high titers against staphylococcal fibrinogen-binding proteins, Ser-Asp dipeptide repeat G (SdrG) and clumping factor A (ClfA), to prevent invasive S. aureus infections in neonates.98 Despite preliminary promising clinical results, Veronate failed to reduce the incidence of neonatal latestage staphylococcal sepsis in phase III.99 It has been suggested that this failure might have been due to the fact that Veronate consisted of a human total IgG preparation and that the ClfA specific IgG concentration may have not been potent enough to neutralize the detrimental effects of ClfA. Alternatively, the ClfA binding antibodies in Veronate were possibly not functional,³⁵ a point which will be addressed further below.

MntC transports essential nutrients during infection. The fourth antigen in the Pfizer S. aureus vaccine is the Manganese Transporter C (MntC), a highly conserved (> 98% sequence identity) lipoprotein that is the surface-exposed metal binding subunit of MntABC, a heterotrimeric membrane transporter responsible for the acquisition of manganese. A primary host defense mechanism against bacterial invasion is the sequestration of metal ions that are essential for bacterial survival.¹⁰⁰ S. aureus and other bacteria have developed approaches to rapidly scavenge divalent cations like manganese and iron from the host when the bacterium establishes infection. As a cofactor for a number of diverse enzymes, manganese plays important roles in bacterial metabolism, cell wall synthesis and virulence.^{101,102} Most notably, it is the sole cofactor for superoxide dismutases, which neutralize superoxide radicals generated during the oxidative burst in the phagosome of macrophages and neutrophils.¹⁰³ S. aureus strains which lack functional MntC display increased sensitivity to superoxide radicals.¹⁰⁴ Therefore antibodies that target MntC have the potential to interfere with two distinct S. aureus virulence mechanisms: nutrient acquisition and phagosome survival.

Pfizer first identified MntC as a potential vaccine target after it was observed that the analogous *S. epidermidis* protein (SitC) was only expressed in vivo or in the presence of human serum.¹⁰⁵ SitC showed > 75% amino acid similarity to MntC and was efficacious in an *S. epidermidis* bacteremia model.¹⁰⁵ In vivo expression analysis of bacteria harvested from the blood at various times revealed that MntC is rapidly upregulated by *S. aureus* in a murine model of infection (**Table 1**). Expression profiles were compared with those for another cation-sequestering protein, IsdB. MntC was found to be consistently expressed earlier during infection in vivo in comparison to other antigens including the IsdB antigen (**Table 1**). Thus, MntC is an important antigen to include in a multicomponent vaccine as it is expressed earlier than many of the surface adhesion molecules.

Preclinical studies demonstrated that MntC could induce a protective immune response.¹⁰⁶ Active immunization studies in a *S. aureus* bacteremia model of infection showed that MntC was effective in reducing the bacterial load of *S. aureus* in the blood. The same antigen also protected against *S. epidermidis* infection (Table 2).¹⁰⁶ This is the first example of a protein antigen that

Antigen	Time [hr]	PFESA0003	PFESA0210	PFESA0182	PFESA0101	PFESA0186	PFESA0155	PFESA0158	PFESA0140	PFESA0266	PFESA0212
		CP Type 8					CP Type 5				
MntC	TO	-	-	-	-	-	-	+	-	-	+
	T1	+	+	-	+	+	+	-	+	+	-
	T4	+	+	+	+	+	+	+	+	+	+
	T6	+	-	+	-	+	+	+	+	+	+
IsdB	TO	-	-	-	-	-	-	-	-	-	-
	T1	-	+	-	-	-	-	-	-	-	-
	T4	-	-	-	-	-	-	+	-	-	-
	T6	+	+	+	+	+	+	+	+	+	-

Table 1. In vivo surface expression of ion scavenging proteins in S. aureus clinical isolates

Immunofluorescence microscopy was performed on *S. aureus* strains harvested from CD1 mice (n = 3) which had been infected by intraperitoneal injection of approximately 2×10^8 colony-forming units at the indicated times after infection using primary rabbit antibodies and an ALEXA488-conjugated goat- α -rabbit antibody.

Table 2. Immunization with MntC reduces the S. aureus bacterial load in a Murine Bacteremia Model (adapted with permission from ref. 106)

Challenge organism	Meta-analysis	Number of animals	Log CFU reduction	P value	
S. aureus	10 experiments	100	0.955	< 0.0001	
S. epidermidis	2 experiments	20	0.95	0.0013	

Groups of 10 CD-1 mice were vaccinated at weeks 0, 3 and 6 with 10 μ g MntC and 22 μ g AlPO4, followed by intraperitoneal challenge on week 8 with 5 × 10⁸ CFU S. aureus Reynolds or *S. epidermidis* O-47. Animals were exsanguinated 3 h after challenge, and serial dilutions of blood were plated to enumerate recovered bacteria. Statistical significance was determined via the Student t test, and a P value of \leq .05 was considered significant.

has the potential to provide protection against both *S. aureus* and *S. epidermidis.* In addition, anti-MntC monoclonal antibodies were identified that could bind both *S. aureus* and *S. epidermidis* cells and were protective in an infant rat passive protection model (Fig. 2). These monoclonal antibodies can be used in serological assays to measure antibodies generated against MntC that compete for binding with the monoclonal antibody to MntC, thereby providing a semifunctional assessment of the potency of antibodies elicited by the MntC vaccine.

Other S. aureus antigens. *Toxins*. A number of *S. aureus* vaccines composed of inactivated toxins or their subunits have been evaluated preclinically.^{35,63} Due to the heterogeneous nature of toxins produced by *S. aureus*, and the fact that toxins are usually released after an infection has been already established, the value of adding toxin components into multicomponent prophylactic vaccine formulation is questionable.

Protein A. Protein A is a highly conserved *S. aureus* cell wall protein encoded by the *spa* gene that binds with high affinity to the Fc region of human IgG1 and IgG2 and mouse IgG2a and IgG2b, thereby inhibiting functional antibodies from blocking key functions such as opsonophagocytoss and adhesion. In addition, Protein A binds to $V_{\rm H}3$ IgM on the surface of B-cell

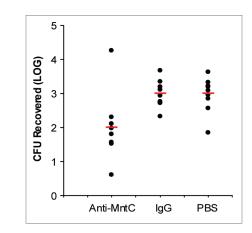


Figure 2. Reduction of *S. aureus* Bacteremia in Infant Rats By Passive Administration of an Anti-MntC Monoclonal Antibody (adapted with permission from Table 4 in ref. 106). Groups of Sprague-Dawley infant rats were immunized intraperitoneally with 0.4 mg of either test mAb or isotype control mAb. Sixteen hours after immunization, rats were challenged intraperitoneally with 1×10^8 colony forming unites of a *S. aureus* clinical isolate, PFESA0140. Four hours later, blood was collected, and serial dilutions were plated to enumerate recovered bacteria. Statistical significance was determined via the Student t-test, and a *P* value of ≤ 0.05 was considered significant.

receptors, resulting in reduced antibody production.¹⁸ Many experts have raised concerns about the possibility that protein A and related proteins, such as Sbi, could render vaccination futile.^{18,65,67} However, it is important to keep in mind that protein A appears to be most relevant for abscess formation and has less impact on the early stages of *S. aureus* infection and dissemination during which time it is not expressed in vivo.¹⁹ Therefore its inclusion in a multi-antigen vaccine designed to prevent infection in the first place may not be appropriate, although it could be considered for therapeutic vaccines.

Suitable antigen expression. Another key feature of a successful vaccine that may not have been sufficiently demonstrated for previous *S. aureus* candidate vaccines is evidence that

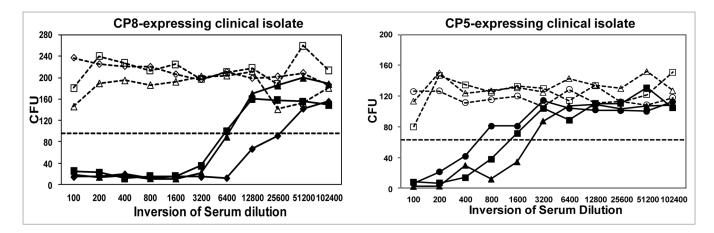


Figure 3. Immune sera from CP-conjugates induce opsonophagocytosis. Sera from three different cynomolgous monkeys vaccinated with either CP8 or CP5 conjugates (each shown with a different symbol) were evaluated for their ability to promote opsonophagocytic killing as described in reference 82. Preimmune sera is shown in dashed lines and post-immune sera, harvested at six weeks post dose, is shown in solid lines. The titer is defined as the reciprocal serum dilution that kills 50% of the test bacteria. The CFU associated with 50% bacterial killing is indicated by the dotted line.

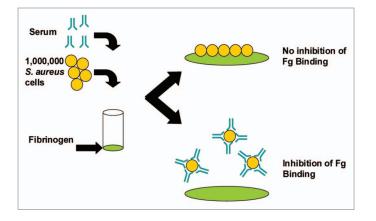


Figure 4. Schematic representation of the S. aureus fibrinogen binding inhibition assay. Microplate wells were coated with fibrinogen (Fg), incubated with blocking solution to prevent non-specific binding and rinsed. Live *S. aureus* cells (1×10^6 colony forming units) anti-sera were mixed and added to the plate. After 30 min incubation at 37°C adherent cells were washed and quantified using the Luciferase-based BacTiter-Glo^{*} reagent, which measures ATP of live bacteria and associated luminescence.

target antigens are expressed to sufficient levels during infection.^{64,67,72} Although the genetic factors influencing the expression of *S. aureus* virulence factors during infection have been described in depth,¹¹⁷ relatively few studies have directly measured antigen expression in vivo. Recent studies using immunofluorescence microscopy in both bacteremia and wound models of infections have been conducted to fully characterize the in vivo expression patterns of capsule and ClfA.¹⁰⁷ It was demonstrated that the expression of these antigens can be temporally different, and is dependent on both the challenge strains examined and the microenvironment encountered during infection. The results also showed that antibody access to ClfA is not impeded by capsule expression.

Demonstration of S. aureus killing and virulence inhibition by vaccine-elicited immune sera. A critical feature of a successful vaccine, which experts believe may not have been suitably addressed in previous S. aureus vaccine clinical trials, is the demonstration that vaccine-elicited antibodies are functional and have sufficient activity and avidity to support opsonophagocytic killing and/or neutralization of important virulence factors.^{108,109} Vaccination with CP5 or CP8 CRM₁₉₇ conjugates elicit robust OPA responses in non-human primates (Fig. 3) and in humans.¹¹⁰ Furthermore, anti-ClfA antibodies generated after immunization with a ClfA-containing vaccine inhibited the binding of S. aureus to fibrinogen in a Fibrinogen Binding Inhibition Assay (FBI assay,¹¹¹ adapted from ref. 86). The FBI assay monitors the inhibition of live S. aureus cells binding to fibrinogen as shown schematically in Figure 4. Such an assay provides a useful readout regarding the effectiveness of a ClfA vaccine component to inhibit this important virulence mechanism. The FBI assay was tested for specificity using a number of S. aureus anti-ClfA monoclonal antibodies (mAbs) that bind with high affinity to native ClfA on the surface of S. aureus. Figure 5 shows the data for a subset of four mAbs, including mAb12-9, the only monoclonal

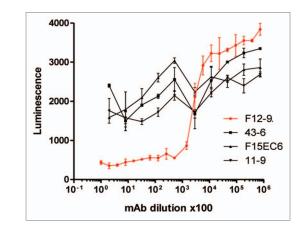
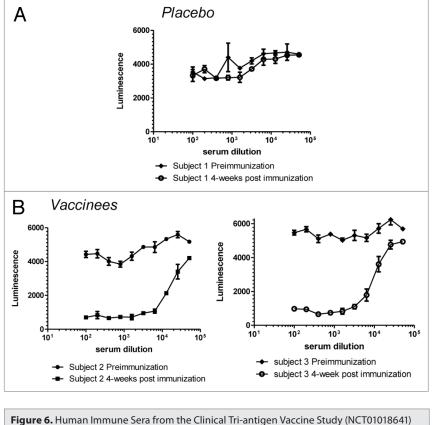


Figure 5. *S. aureus* FBI assay is specific for anti-ClfA functional monoclonal antibodies. Serial dilutions of monoclonal antibodies (mAb) were pre-incubated with *S. aureus* strain PFESA0237 and their binding to immobilized fibrinogen was assessed as described in the legend to **Figure 4.**¹¹¹ Despite similar binding affinities,only the functional mAb 12–9 shows binding inhibition in a dilution-dependent manner.

antibody that was protective in an animal model of staphylococcal infection and, importantly, the only antibody in this subset that could inhibit the binding of *S. aureus* to the fibrinogen. Despite similar binding affinities, only mAb 12–9 shows binding inhibition in a dilution-dependent manner. These results (1) further confirm the important role of ClfA in virulence and (2) underscore the important distinction between binding affinity and functional activity.

Clinical sera from the Pfizer SA3Ag vaccine study (described below) were evaluated using the exploratory FBI assay. No FBI activity was detected in unvaccinated subjects and complete inhibition of Fg binding was observed for most vaccinated subjects (28/32) (Fig. 6). It is important to note that, despite the frequency of *S. aureus* colonization and the presence of pre-existing antibodies, most individuals do not have circulating functional antibodies to this important virulence factor. Encouragingly, immunization with a ClfA-containing vaccine does induce high titers of such functional antibody, illustrating the critical importance of demonstrating the functionality of antibodies generated during vaccine development.

Current status in the clinic. As described on www.clinicaltrials.gov, Pfizer recently completed a phase I trial to evaluate a trivalent vaccine composed of two capsular polysaccharide (CP) conjugates and Clumping Factor A (CP5, CP8 and rmClfA) in healthy adults (18 to 85 y of age) (ID#NCT01018641) and currently has phase I/II trials in progress to evaluate three ascending doses of a tetravalent vaccine (consisting of CP5, CP8, rmClfA and MntC) (ID#NCT01364571 and ID#NCT01643941) in the same target population. Although CPs are produced by bacteria to be antiphagocytic, Pfizer and others have shown in *S. aureus*, anti-CP antibodies mediate killing by opsonophagocytosis. ClfA is associated with bloodstream infections and facilitates the binding of *S. aureus* to fibrinogen.⁸⁹ The vaccine in the initial study was reported to be well-tolerated and able to elicit robust functional antibodies after a single dose.¹¹⁰ Of significance was



Prevents *S. aureus* (strain PFESA0237) from Binding Fibrinogen. Serial dilutions of serum taken from human subjects either before or four weeks after immunization with either a placebo (**A**) or the trivalent vaccine (**B**) were tested for activity in the FBI assay as described in the legend to **Figure 4**.

the use of the FBI assay described above to demonstrate that healthy humans do not have functional antibodies that can prevent deployment of this virulence mechanism. In contrast, high inhibitory titers were observed after vaccination.¹¹¹

Other Considerations

Clinical trial design. Several recent reviews have included detailed discussions on key considerations for the design of a successful clinical trial for *S. aureus* vaccines.^{35,64,66} In addition to the relevant, robust functional assays to characterize initial immune responses and a careful selection of patient populations that would be most appropriate for initial proof-of-concept efficacy evaluations, the definition and timing of the disease endpoint is also critical.

Risk factors. It is almost certain that selection of the appropriate target population for a *S. aureus* vaccine is a key determinant for the successful demonstration of its efficacy. The risk factors for invasive *S. aureus* infection are well-defined for both community and health-care settings⁶⁶ and notably distinct from each other, and with the former consisting of chronically immunocompromised patients and the latter consisting of patients in the ICU or undergoing surgery or recipients of foreign bodies

such as catheters, implants or other medical devices.^{112,113} It is therefore implicit that vaccines targeting these distinct populations would differ in their development strategies.⁶⁴

HA-MRSA vs. CA-MRSA. Strains causing infections differ between hospital and community. Methicillin-resistant *S. aureus* (MRSA), historically associated with nosocomial infections, is now highly prevalent in both healthcare and community settings.^{114,115} However, MRSA strains causing disease in the community are not only phylogenetically and phenotypically distinct from those found in the hospital,¹¹⁶ but also differ in the degree²⁷ and site¹¹⁷ of colonization and often demonstrate different clinical disease manifestations.¹¹⁸ These differences must also be taken into account during the design of an effective vaccine.

Conclusion

S. aureus is a challenging vaccine target due to complexity of its pathogenesis involving a multitude of virulence factors. However, these pathogenic mechanisms can be addressed by eliciting functional antibodies to an appropriate selection of *S. aureus* antigens that are expressed in vivo early during the infection cycle. Pfizer's *S. aureus* investigational vaccine has been designed to address three major virulence mechanisms associated with staphylococcal disease: immune evasion (capsule), binding

to host cells and immune evasion (ClfA) and nutrient acquisition (MntC). The need to generate functional antibodies that are capable of mediating measurable opsonophagocytic killing has been addressed by inclusion of conjugated capsular polysaccharide antigens CP5 and CP8. We have demonstrated that anti-ClfA is an important virulence factor and that vaccine-elicited antibodies can completely inhibit S. aureus binding to human fibrinogen which is required to prevent ClfA mediated pathology. We have also shown that MntC is protective in preclinical models of infection and others have demonstrated that S. aureus lacking MntC are more susceptible to oxidative stress. Unlike monovalent vaccine candidates described in the past that were not successful in preventing S. aureus infections, it is hoped that inclusion of multiple S. aureus antigen targets involved in several pathogen defense mechanisms and that are expressed early in the infectious cycle, will be sufficient in preventing infection and disease with this important pathogen.

Disclosure of Potential Conflicts of Interest

All authors are employees of Pfizer and as such were paid to do this work and may have shares in the company. ASA was a previously employed by Merck and may have shares in the company.

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