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Integrating complex host-pathogen immune environments into *S. aureus* vaccine studies

Chih-Ming Tsai^{1,#}, Irshad A. Hajam^{1,#}, JR Caldera^{1,2,#}, George Y. Liu^{1,3,*}

¹Division of Infectious Diseases, Department of Pediatrics, University of California San Diego, La Jolla, CA 92093, USA.

²Department of Biomedical Sciences, Cedars-Sinai Medical Center, Los Angeles, CA 90048, USA.

³Division of Infectious Diseases, Rady Children's Hospital, San Diego, CA 92123, USA.

Abstract

Staphylococcus aureus (SA) is a leading cause of bacterial infection and antibiotic resistance globally. Therefore, development of an effective vaccine has been a major goal of the SA field for the past decades. With the wealth of understanding of pathogenesis, the failure of all SA vaccine trials has been a surprise. We argue that experimental SA vaccines have not worked because vaccines have been studied in naïve laboratory animals, whereas clinical vaccine efficacy is tested in immune environments reprogrammed by SA. Here, we review the failed SA vaccines that have seemingly defied all principles of vaccinology. We describe major SA evasion strategies and suggest that they reshape the immune environment in a way that makes vaccines prone to failures. We propose that appropriate integration of concepts of host-pathogen interaction into vaccine study designs could lead to insight critical for the development of an effective SA vaccine.

Graphical Abstract

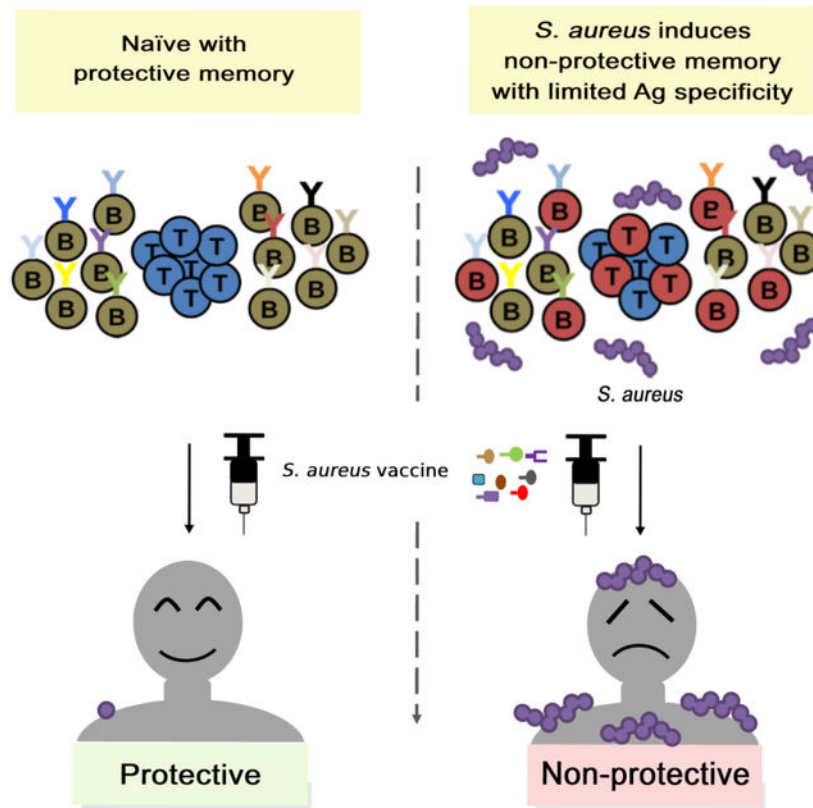
*Correspondence: gyliu@health.ucsd.edu.

#These authors contributed equally to the review.

Declaration of Interests

The authors declare no conflict of interest

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The failure of all *S. aureus* vaccine trials has been a conundrum. Tsai et al. propose that staphylococcal vaccine failures result from underappreciation of how the pathogen preprograms the host immune system for vaccine failures. Understanding *S. aureus* modulation of vaccine responses may hold the key to developing successful vaccines.

Keywords

Staphylococcus aureus; vaccine; original antigenic sin; evasion mechanisms; T cells; B cells; antibodies; pathogenesis

Introduction

Staphylococcus aureus (SA) is a versatile pathobiont that is exceptionally well-adapted to co-exist with the human host. The Centers for Disease Control estimated that 120,000 cases of SA bloodstream infections and 20,000 associated deaths occurred in the United States in 2017 (Kourtis et al., 2019). Persistence of methicillin-resistant SA (MRSA) infections in hospital and community settings has increased the use of once restricted antibiotics and led to inevitable acceleration and spread of antibiotic resistance (Chambers and Deleo, 2009). The importance of SA to human health as well as its versatility as a pathogen have drawn abundant research interest. As a result, much knowledge has been garnered on how SA interacts with the host in experimental models. In comparison, translational SA research has lagged even though differences in SA interaction with humans and mice have

been appreciated for a long time. When called upon to address human relevant issues in the past two decades, the field has repeatedly failed to meet the challenge. For example, mouse studies of Panton-Valentine Leukocidin elaborated by Community-Associated MRSA proved to be unhelpful, and the pathogenic role of the toxin was debated for a decade until it was determined that rodents are not the best tool to study the human-tropic toxin (Spaan et al., 2013). In response to calls for a vaccine to control the expanding MRSA crisis, more than twenty human active and passive vaccine trials have been conducted (Armentrout et al., 2020; Miller et al., 2020). Yet, to date, an effective vaccine remains elusive. Although diverse opinions remain on why the vaccines failed, experts have come closer to an agreement that small tweaks in vaccine designs are unlikely to solve the vaccine conundrum.

It is noteworthy that most SA vaccine studies have been carried out in naïve laboratory animals. The commonly used mouse models show little evidence of prior human SA interaction, apart from the occasional encounter with murine SA that do not produce many of the virulence factors expressed by the human-tropic SA (Holtfreter et al., 2013). In comparison, humans are exposed to SA during the first months of life and show evidence of continued exposure with increasing anti-SA titers into adulthood (Lebon et al., 2008; Li et al., 2021). SA uses diverse strategies to evade both innate and adaptive immune responses and maintain coexistence with the human host. Many experts have pointed to these mechanisms as likely reasons for the vaccine failures, but direct evidence is lacking. However, there is growing momentum to seek out more fundamental reasons for the failure of the vaccines (Teymournejad and Montgomery, 2021). In this review, we argue that a translational approach that fully integrates the prior SA experience of the host with testing of vaccine efficacy is an important step towards building that translational bridge. We provide an overview of the unsuccessful SA vaccine trials and discuss the inability of traditional vaccine approaches to solve the SA vaccine conundrum. We review the primary SA T and B cell evasion mechanisms that conceivably could alter vaccine efficacy by altering the existing immune environment. We then reimagine what vaccine response would look like in that environment, drawing on the concept of original antigenic sin that has been applied to viral pathogens to explain vaccine failures, but underlining the fine differences that exist with immune imprinting as it applies to SA and viral pathogens. Due to space limitation we apologize to many contributors in the field whose important work are not adequately presented.

S. aureus vaccine failures

SA has assumed notoriety as the leading cause of bacterial infections with the emergence of MRSA, first in intensive care units within healthcare settings in the 1980s, and subsequently in communities (Chambers and Deleo, 2009). SA infections range from the common soft tissues infections to invasive diseases that carry significant mortality even with appropriate antibiotic treatment. Increased MRSA burden over the past decades have also broadened the use of second and last line antibiotics, thus driving antibiotics such as vancomycin to the edge of obsolescence (Chambers and Deleo, 2009). Combined, these events have led to the consensus on the urgency of developing a SA vaccine in the Threats Report developed by the CDC (2019).

Following the successes of capsular polysaccharide vaccines against *H. influenzae* and *S. pneumoniae* and prompted by promising pre-clinical vaccine studies against SA capsular polysaccharides (Yoshida et al., 1987; Lee et al., 1988; Fattom et al., 1996), early phase II and III SA vaccine trials targeted the Type 5 and Type 8 staphylococcal capsular polysaccharides (Fattom et al., 2015; Rupp et al., 2007; Robbins et al., 2004). Despite demonstrating a trend towards protection at the 40-week timepoint, neither the one-dose regimen in the first trial, nor two-dose approach in the second trial provided protective efficacy in end-stage renal disease hemodialysis patients (Fattom et al., 2015). Among suggested reasons are low capsule expression during human infections and interference with capsular binding by another antibody to surface polysaccharide PNAG through idiotypic binding (Skurnik et al., 2010). Additionally, the target patient population could have decreased complement and phagocyte functions and might not have been an optimal population to study for an initial vaccine trial (Fattom et al., 2015).

With the unsuccessful anti-capsular vaccine approach, several of the next trials targeted cell-wall anchored proteins, in part, motivated by the success of a large number of experimental vaccines against surface antigens (Stranger-Jones et al., 2006). Unexpectedly, none of the trials were successful. The most glaring of the failures was the Merck V710 double-blinded randomized vaccine trial for the prevention of SA infection after cardiothoracic surgery (Fowler et al., 2013). The vaccine that targeted the iron-regulated surface determinant protein B (IsdB) showed significant promise in several murine models of SA infection (Brown et al., 2009; Kuklin et al., 2006; Stranger-Jones et al., 2006). Notably, beyond failing to reduce the incidence of SA in vaccinated subjects, those who were infected with SA after receiving the vaccine had a five-fold increase in mortality (Fowler et al., 2013). Almost all vaccine recipients who died of SA infection had undetectable levels of serum IL2 and IL17A prior to vaccination, suggesting the potential contribution of specific host factors to infection severity (McNeely et al., 2014). A subsequent phase IIB vaccine trial targeted 4 SA surface antigens (MntC, ClfA, CPS5 and 8), based on the concept that immunizing against multiple antigens would be more efficient, but the trial was stopped despite inducing persistent and robust titers against all the antigens over 36 months because of low probability that efficacy objectives, protection against infection over 180 days from time of surgery, would be achieved (Gurtman et al., 2019).

With the failure of vaccines targeting cell surface antigens, efforts turned to neutralization of toxins that are direct source of immunopathology. It was argued that while this approach would not directly eliminate SA, anti-toxin vaccine strategy has been successful against pathogens such as diphtheria, tetanus, and pertussis. One ongoing trial targets seven toxins, including α -toxin, PVL, SEA, SEB, and TSST-1 (Aman, 2018), with several other trials concurrently targeting α -toxin and other cell surface proteins (Miller et al., 2020).

Since active vaccination is a multi-step process involving collaboration between many immune cell types, human and mouse differences at any step have the potential to lead to differences in vaccine efficacy. Hence, passive application of monoclonal antibodies that have been preselected for affinity and antigen neutralization could bypass these potential pitfalls. To date, SA vaccine trials have tested efficacy of monoclonal antibodies against a variety of SA antigens, including toxins (α -toxin), PAMPs (LTA) and surface antigens (ClfA

and CP5/CP8), in prophylactic or treatment settings (Francois et al., 2021; Rupp et al., 2007; Weems et al., 2006; Weisman et al., 2011; Yu et al., 2017). Disappointingly, once again, none of the trials achieved their efficiency targets. In the case of the anti- α -toxin monoclonal antibody Suvratoxumab, application of a 5-gram dose that achieved neutralizing activity of 156 IU/ml on day 2 and 33 IU/ml at 90 days failed to significantly reduce the incidence of pneumonia in ICU ventilated patients at 30 days (Francois et al., 2021; Wu et al., 2018), although promising sub-analysis of patients less than 65 years of age has led to a planned phase III trial.

A different vaccine approach aimed at promoting T effector functions to control SA diseases. Th1 and Th17 immunity appears to be important for protection against a variety of SA infections (Brown et al., 2015; Lee et al., 2020; Paterson et al., 2020). One promising experimental vaccine promoting Th1/Th17 immune response was a vaccine that targeted candida antigen Als3p-N that bears structural similarity to SA ClfA (Lin et al., 2009). The vaccine advanced to a phase II trial that aimed to prevent SA nasal colonization among military recruits (Schmidt et al., 2012). Results of the trial are currently pending.

Overall, repeated failures of the human trials have eroded confidence of the field in traditional vaccine approaches and have fostered a growing consensus that more fundamental understanding of SA vaccination in humans and animals is needed to overcome the current impasse in vaccine development. SA's versatility as a pathogen and its commensal relationship with the human host likely need to be accounted for in rethinking SA vaccine approaches.

Clinical evidence of protective humoral and cell mediated immunity

Human SA colonization occurs from early childhood (Lebon et al., 2008). Between colonization and occasional infections, children develop levels of SA-specific antibodies and T cells that increase with the age (Li et al., 2021). Despite that, only moderate level of immunity develops. Studies have shown that individuals chronically colonized with SA are more frequently infected, but the severity of invasive SA disease if that occurs is reduced compared to non-colonizers, suggesting some level of acquired protection (Wertheim et al., 2004). Experiments in mice with skin or bloodstream reinfection, to a large extent, reflect the human observation, although Major Histocompatibility Complex (MHC)-restriction of the mouse strain appears to influence protection to reinfection (Si et al., 2020).

When further dissected, the role of B cells in protection is thought to be limited, since individuals with B cells deficiency are not more susceptible to SA infections compared to normal individuals (Fowler and Proctor, 2014). However, there is support for the protective role of some toxin-specific antibodies against select SA diseases. For example, individuals with staphylococcal toxic shock syndrome (TSS) have significantly lower titers of antibody to TSS Toxin (TSST) (Bonventre et al., 1984), which is the rationale behind the use of IVIG, pooled immunoglobulins obtained from at least one thousand donors, in the treatment of patients with TSS (Darenberg et al., 2004). Likewise, convalescent antibodies to exotoxin α -toxin in patients with SA infection correlate with protection against subsequent SA infections (Fritz et al., 2013), while higher antibody titers against several

secreted toxins collectively correlate with protection against sepsis syndrome in patients with SA bacteremia (Adhikari et al., 2012). In a model of SA-mediated septic arthritis, B-cell-deficient agammaglobulinemic mice did not develop more severe disease or increased bacterial burden compared to wild-type mice (Gjertsson et al., 2000). Similarly, experiments using Rag2-deficient mice demonstrated that, while adaptive immune cells are activated during SA infection and are required for prolific inflammatory cytokine secretion, the absence of mature B- and T-cells only had a local and temporal effect on bacterial clearance in the liver during early sepsis and had no significant effect altogether in late sepsis.

In contrast to B cells, T cells appear to have a more substantial role in containing SA infections in humans and experimental mouse models (Cho et al., 2010; Levy et al., 2016; Miller and Cho, 2011). Robust evidence supports the central role of CD4⁺T cells, particularly Th1 and Th17 cells, in mediating protection against SA infections (Brown et al., 2015; Lee et al., 2020; Paterson et al., 2020). Notably, the importance of Th17 cells can be appreciated in individuals with STAT3 loss-of-function mutations or hyper-IgE syndrome that interferes with the differentiation of Th17 cells (Ma et al., 2008; Milner et al., 2008), which thus leads to recurrent and severe mucocutaneous SA infections. Likewise, HIV-positive patients who experience significant depletion of Th17 cells early in the course of their HIV infection are poor controllers of SA skin and soft tissue infections (Brenchley et al., 2008; El Hed et al., 2010; Hidron et al., 2010). Lastly, patients with atopic dermatitis have increased skin colonization and superinfection with SA, with decreased IL-17 pathway cytokines and increased Th2 cytokines, including IL-4 and IL-13, in lesional skin (Guttman-Yassky et al., 2008). Along with Th17, IFN- γ produced by Th1 cells has also been implicated in protection against SA skin and bloodstream infections (Beekhuizen and van de Gevel, 2007; Brown et al., 2015). Another CD3⁺T cell, which displays neither CD4 nor CD8 coreceptor on its surface, is the gamma-delta T cell that has been reported to mediate protection against skin and soft-tissue SA infections in an IL-17 dependent manner (Dillen et al., 2018; Leyva-Castillo et al., 2021; Marchitto et al., 2019). The specific roles of each subset of CD4⁺T cell during SA infections is reviewed elsewhere (Armentrout et al., 2020) and is beyond the scope of this review.

S. aureus manipulation of humoral responses

SA evades host humoral defenses through factors that subvert antibody functions and appropriate B-cell development (Figure 1). Principally, the well-characterized SA protein A (SpA) antagonizes humoral immunity through interactions with both the antibody constant fragment (Fc), and the B-cell receptor (BCR) variable domain. Early reports demonstrated that SpA non-specifically binds human gamma-globulins at the Fc receptor to inhibit SA phagocytosis and bacterial killing (Forsgren and Quie, 1974; Forsgren and Sjoquist, 1966). It was later elaborated that SpA binding of antibody Fc fragments can also result in the formation of multi-molecular complexes that mediate intracellular SA survival and lead to systemic dissemination of surgical site infection (Nishitani et al., 2020). This IgG-binding function of SpA is mirrored by a similar virulence factor, Staphylococcal immunoglobulin-binding protein, Sbi, to further expand SA's capacity to evade antibody-mediated clearance (Zhang et al., 1999; Zhang et al., 1998).

In addition to SpA's impact on antibody-dependent opsonophagocytosis, Goodyear and Silverman provided evidence of SpA's superantigenic function by tracking the fate of SpA binding peripheral B-cells and demonstrated that this association promotes supraclonal deletion by apoptosis. By crosslinking with a conserved V_H region of the BCR variable domain, SpA induces a B-cell activation state that is driven towards programmed cell death (Goodyear and Silverman, 2003). Consistent with the detrimental effects of SpA, mice and guinea pigs infected with a functionally SpA-deficient mutant show improved phagocytosis of SA and mount a protective B-cell response against lethal SA challenge (Falugi et al., 2013; Fattom et al., 2015).

Amidst its deleterious effects, SpA can also trigger vigorous B-cell proliferation. By cross-linking BCRs, SpA sensitizes B-cells for the recognition of TLR2 ligands to promote expansion of intracellular IgM-expressing B-cells. Further investigation, however, revealed that these B-cells fail to induce significant secretion of the SA-targeting immunoglobulins (Bekeredjian-Ding et al., 2007). SpA-mediated cell activation selectively triggers the expansion of IL-10-secreting regulatory B-cell subsets. In cooperation with plasmacytoid dendritic cells, SpA interaction with B-cells strengthens the characteristic immunosuppressive IL-10 response associated with SA infections (Parcina et al., 2013). Thus, whether by clonal deletion or non-productive cellular expansion, the impact of SpA on B-cells allows SA to efficiently evade humoral immunity. As further evidence for these mechanisms, Schneewind and Missiakas developed a nontoxicogenic SpA vaccine that overcomes SpA's dual functions by antagonizing the effects of SpA on both antibodies and B-cells. Vaccination with SpA_{KKAA}, a variant that does not bind Fc_γ or Fab V_H3, promoted opsonophagocytic clearance, as well as a more robust antibody response against many SA antigens in mice (Kim et al., 2010).

Many studies have examined the implication of these findings in humans by assessing the presence and efficacy of anti-SA antibodies naturally circulating in human serum and induced after SA infection. A study of plasmablasts from SA-infected subjects showed a focused immunodominant response to SpA and a more limited response to other SA virulence factors, consistent with the proposed superantigenic mechanisms of SpA (Pauli et al., 2014). However, antibody profiling studies from other groups have shown the near ubiquity and high abundance of anti-SA antibodies in the healthy population, which further increase during SA infections (Dryla et al., 2005; Radke et al., 2018; Romero Pastrana et al., 2018). Addressing the functionality of these antibodies, we showed that adoptive transfer of sera from healthy children are largely non-protective in a murine SA challenge model. In comparison, forty percent of convalescent serum samples from children with invasive SA disease are able to reduce SA burden at 4–6 weeks post-infection, but not at 6 months (Tsai et al., 2021). These data are consistent with the relatively non-protective role of humoral immunity in humans, particularly in children, despite abundant antibody production. It remains unclear how SA manages to make most SA-specific antibodies non-protective.

In addition to SpA, a recent study showed that SA leukocidins also play a role in modulating host humoral responses (Tam et al., 2020). The authors showed that infection of mice with a *lukED hlgACB* double mutant SA resulted in increased anti-SA antibody levels

compared to infection with the isogenic WT SA. Hence targeting of SA leukocidins could improve efficacy of vaccines.

S. aureus manipulation of T cell responses

SA targets effector functions of CD4⁺T cells mainly with two classes of virulence factors: superantigens and toxins (Goldmann and Medina, 2018; Thammavongsa et al., 2015; Xu and McCormick, 2012) (Figure 2). Superantigens like TSST-1 and staphylococcal enterotoxin B bind class II MHC molecules and to the variable region of a specific V β chain of the TCR to induce potent activation of about twenty percent of all peripheral T cells (Xu and McCormick, 2012). This non-specific superantigen-mediated T cell activation prevents the development of a focused and coordinated immune response and leads to the loss of overall receptor diversity and lack of antigen-specific protective T cell responses.

SA additionally produces an array of functionally diverse toxins, including leukocidins, hemolysins, and phenol soluble modulins (PSMs), which launch a fierce attack on the host immune system to subvert protective T cell responses (Berends et al., 2019; Richardson et al., 2018; Spaan et al., 2017). Leukocidins are secreted factors that specifically target human and mouse lymphocytes. For example, LukED targets and kills CCR5-positive expressing Th1 and Th17 cells (Alonzo et al., 2013). Similarly, α -toxin induces programmed cell death of human and mouse IFN- γ expressing T cells during MRSA infection (Bonifacius et al., 2020; Nygaard et al., 2012). The detrimental effect of α -toxin on memory T cells is evidenced by the development of enhanced specific memory T cell response in mice born to Hla_{H35L}-immunized dams and subsequently challenged with SA (Lee et al., 2020).

In addition to toxin-mediated killing of T cells, SA leukotoxins LukAB and LukED, induce direct killing of human dendritic cells (DCs) (Alonzo et al., 2013; Berends et al., 2019), which are a central player in priming of adaptive immune responses. PSMs also disturb the adaptive immune response via the induction of tolerogenic dendritic cells (DCs) (Richardson et al., 2018; Schreiner et al., 2013). PSM-treated DCs produce high level of IL-10 and increase the frequency of FOXP3⁺ regulatory T cells, which have been shown to suppress both Th1 and Th17 cellular immune responses (Mondal et al., 2012; Schreiner et al., 2013). Furthermore, SA infection expands the myeloid-derived suppressor cell (MDSC) population, which are well-known to suppress effector T cell functions through the secretion of IL-10 (Heim et al., 2015; Peng et al., 2017). Consistently, a report showed that SA peptidoglycan-induced IL-10 prevents Th1 and Th17 cellular immune responses (Frodermann et al., 2011). This finding was recently corroborated by a study from our group demonstrating more specifically that O-acetylation of peptidoglycan suppresses Th17 cell responses in an IL-10 dependent manner, and that compared to wild-type mice, IL-10 deficient mice immunized with live SA vaccine had improved Th17 response, and thus, bacterial clearance upon challenge (Sanchez et al., 2017). Adoptive transfer of CD4⁺T cells from the immunized IL-10 deficient mice into naïve recipient mice conferred significant protection against SA infection. These results indicate that IL-10 produced during primary SA infection is likely one of the unique and important immune evasion strategies employed by SA that contributes to the lack of protective memory CD4⁺ T cell responses. Further elucidating the mechanism by which IL-10 subverts CD4⁺T cell immunity, a study in a *Mycobacterium tuberculosis*

(Mtb) model showed that CD4⁺ T cells primed in an IL-10-enriched environment are functionally incompetent and unable to control the infection. This non-protective phenotype was stable and maintained even after the IL-10-modulated T-cells were transferred into IL-10-low recipients (Ferreira et al., 2021). Whether a similar effect occurs with SA-induced IL-10 warrants thorough investigation. In line with the demonstrated effect with Mtb, SA-induced IL-10 could be postulated as a survival strategy exploited by the pathogen, whereby it may help the pathogen persist and thrive within the host while serving to limit inflammatory damage to host tissues and organs.

Yet another mechanism employed by SA to interfere with the development of effective T cell response is the molecular mimicry of host immune components. SA secretes a Class II MHC analog protein, MAP, which has been shown to impede T cell proliferative response and induce Th2 cell differentiation. Pertinently, Th2-associated IL-4 cytokine suppresses IL-17 response (Leyva-Castillo et al., 2021).

SA T cell evasion mechanisms are more difficult to study clinically than antibody responses. Overall, staphylococcal superantigen effect on the induction of T cell anergy has been reported in atopic dermatitis and psoriasis (Yarwood et al., 2000), but it unclear if it has a more than a transient effect on the human T cell repertoire to affect SA vaccination. SA modulation of host cytokine responses (i.e. elevation of IL-4 and IL-10) has also been demonstrated in atopic dermatitis, as well as after systemic infections (Rose et al., 2012). Particularly, the association of mortality with IL10 in SA bacteremia is well documented (Leyva-Castillo et al., 2021; Rose et al., 2012). In a SA study of children, infections, irrespective of invasive or non-invasive nature, correlated with global impairment of anti-SA Th17 responses compared to healthy (colonized) children, suggesting a mechanism whereby infection to SA drives non-protective outcome (Li et al., 2021).

Vaccination in a *S. aureus* reprogrammed host environment

Considering an immune system shaped by the above SA mechanisms, we ask how SA vaccines would perform in such an environment instead of the naïve laboratory mouse environment. Although SA frequently colonizes the human host, it is unlikely that the colonizing SA would interact in a significant way to affect vaccine response directly. Hence, we will not consider direct interaction of the vaccine with SA virulence factors in this discussion.

SA vaccine could be expected do one of two things: It could prime for a new cellular or humoral response, or it could recall an anti-staphylococcal memory response. If the memory response is protective, vaccine would be anticipated to amplify the protective memory response, as is shown when patients who recovered from SARS-CoV-2 infection are vaccinated against SARS-CoV-2 (Stamatatos et al., 2021). Also consistent is the success of vaccine against LukAB, a presumed protective antigen, in SA-colonized minipigs (Fernandez et al., 2021). Conversely, most host antibody responses to SA are presumed to be non-protective (Miller et al., 2020), and vaccine recall of these responses would be expected to be non-protective. Hence, a SA capsular polysaccharide vaccine that was effective in

naïve mice was shown to be not protective in SA-colonized minipigs, consistent with human trial findings (Fernandez et al., 2021; Fattom et al., 2015).

This hypothesized vaccine response draws from the concept of Original Antigenic Sin (OAS) proposed by Thomas Francis, Jr in 1960 (Francis, 1960). It describes the recall of a memory response to a primary influenza infection by secondary viral exposures or vaccinations. Because viral antigenic shift in influenza occurs seasonally, the antigen seen by the immune system subtly differs between exposures. The efficacy of the recalled memory response is determined to a large extent by the antigenic distance between the initial and subsequent encounters. Additionally, the recalled response could reduce *de novo* priming of an effective response against the new antigen. As such, OAS could be protective, non-protective, or even suppressive.

In the case of SA, the antigens seen by the immune system with primary infection and subsequent vaccination could be the same because of the routine practice of selecting conserved targets for immunizations. Because most of the humoral responses are presumed to be non-protective, vaccine reliance on recall of the initial response would suggest that many vaccines to SA would be non-protective. However, various factors could modify this response to alter overall SA vaccine efficacy.

Generation of a protective *de novo* response

Can OAS be overcome? If the recalled memory response is modest, it is conceivable that the SA vaccine could induce a *de novo* response from the naïve pool of T and B cells. It would be intuitive to think that this *de novo* response would be non-protective if the host response to the same antigen is non-protective in the context of infection. Surprisingly available data suggest that effective vaccines are readily made to all types of SA antigens, even those cell-surface antigens that are presumed to be “non-protective” (Stranger-Jones et al., 2006). This observation would thus suggest that the context in which the antigen is presented, either infection or adjuvant, could lead to opposite protective outcomes. To rationalize this outcome, it has been well established that both Fab and Fc domains contribute to protective function of antibodies through their interaction with SA antigens and host immunocytes (Bennett et al., 2019; Chen et al., 2020). Hence, changes, for example, in Fc glycosylation as a result of priming with adjuvant or through infection could conceivably lead to differences in protection. It is less obvious how a protective and a non-protective Fab response could be mounted to the identical antigen. However, there are examples of how pathogens have directed immune responses to immunodominant but non-protective subdomains on the microbial cell-surface, with unmasking of protective epitopes only by selective subdomain vaccines (Novotny and Bakaletz, 2003; Wrightsman et al., 1994). Irrespective, protective antibodies to the “non-protective” antigen could conceivably be made through vaccination, although there are additional factors within the SA reshaped environment that could limit their efficacy as discussed below.

Interference with priming

Hypothetically, recalled memory cells could limit *de novo* T or B cell priming through direct competition for space and cytokines. The extent of interference would depend on

the magnitude of the recalled response, the vaccine antigen concentration, and the antigen presenting cell numbers. Prior colonization or infection with SA could also modify the T and B cell repertoire through depletion or other suppressive mechanisms associated with SA T and B cell superantigens. Silverman described the generation of holes in the B cell repertoire as a result of interaction with SpA antigen. When SpA treatment is stopped, conventional B-2 repertoire normalized, but the “hole” in B-1 repertoire persisted (Silverman et al., 2000). It is less clear to what extent SA superantigens affect T and B cell repertoires in humans.

Pre-existence of non-protective specific antibodies can also interfere with de novo T or B cell priming by two distinct mechanisms (Bergstrom et al., 2017; Getahun and Heyman, 2009). The antibodies could bind the vaccine antigen and facilitate its clearance. Alternatively, antibody binding to the vaccine antigen could mask and thereby dampen de novo priming of naïve T or B cells. Modeling both mechanisms using specific IgG to sheep red blood cells, Heyman and colleagues showed suppression of naïve and memory B and T cell activation via both mechanisms although epitope masking was predominant.

Cytokine modulation of T and B cell development

Assuming that the vaccine antigens successfully initiate priming of potentially protective naïve T or B cells, exposure to appropriate cytokines is still required for the development of effective anti-SA immunity. Elevated IL-4 and IL-10, in association with several types of SA infections, have the potential to undermine development of protective T and B cells (Leyva-Castillo et al., 2021; Rose et al., 2012). If the recalled SA-specific T or B cells turn out to be the primary producer of IL10 or IL4 (Sanchez et al., 2017), then the recall response has the potential to further reduce vaccine efficacy through the cytokines’ suppressive properties. By the same argument, it might be possible to drive T or B cells towards a protective phenotype using Th1/Th17 adjuvants. Plasticity might even allow for the conversion of non-protective memory cells as shown in a study where a protective vaccine Th1/Th17 memory response to SA TSSST-1 is shown to be lost because of memory cell conversion to a IL10 regulatory phenotype (Narita et al., 2019).

Direct antibody competition

Assuming that vaccination is able to induce de novo protective antibodies, efficacy of the vaccine would be determined by the outcome of competition between the protective and recalled non-protective antibody responses. In support of this mechanism interference, a study in a rodent malaria model demonstrated the capacity of a non-neutralizing monoclonal antibody to interfere with a protective antibody in vitro and in vivo even though they bind non-overlapping regions of the same sporozoite antigen. The authors pointed to the data as proof of principle demonstration that pre-existing non-protective antibodies could make malaria vaccines less efficacious for malaria-exposed individuals in endemic areas (Vijayan et al., 2021). Unlike influenza vaccine antigens which recall cross-reactive antibodies that are not expected to bind with similar affinity as the de novo primed antibody response, vaccine- and infection- induced antibodies target the same SA antigen. Hence, the likelihood of interference is greater. This mechanism of vaccine suppression could have significant implications on SA vaccinology because it could explain not only the pervasive failure in

active vaccination, but may also explain the disappointing outcomes in passive immunization platforms.

Post-vaccine effect

Even after vaccination, SA has the potential to further modulate anti-SA effector functions. In a murine model of SA re-infection, Keener et al. showed that SpA alters the fate of plasmablasts and plasma cells by enhancing the short-lived extrafollicular response and reducing the pool of long-lived plasma cells (Keener et al., 2017). The effect of this modulation on antibody production is corroborated by evidence of lower specific antibody titers to several SA antigens after infection with SpA-expressing wild-type compared to a functionally SpA-deficient strain (Falugi et al., 2013). SA also secretes the toxin LukED that specifically targets and kills the predominant CCR5-positive effector memory T cell population (Alonzo et al., 2013). In another mouse study, a mutant TSST-1 vaccine induced a protective specific Th17 response one week after vaccination. However, anti-TSST immunity was lost after 12 weeks and was shown to be related to IL10 secretion by the memory cells which suppresses IL17 production (Narita et al., 2019).

Other human-specific considerations

Microbiome in laboratory animals also has the potential to confound vaccine data as highlighted by a recent discovery that laboratory mice with “wild” microbiome have immune characteristics that are more aligned with human immune responses. The particular effect of microbiome on vaccine-induced immunity was shown in laboratory mice cohoused with pet-store mice (Fiege et al., 2021). These mice showed dampened influenza vaccine-induced humoral responses and poor control of influenza infection compared to the laboratory mice. Additionally, heterosubtypic protective T cell responses were compromised in co-housed mice, indicating the influence of microbiome on vaccine-induced immunity. This has helped to further focus attention on the limited translational potential of current mouse models.

Unrelated to the human environment, SA tropism for human-encoded immune factors is another feature that could drive discrepant results in the murine and human hosts (Spaan et al., 2017). As a human pathogen, SA elaborates many virulence factors, particularly toxins, that interact poorly with the murine host (Spaan et al., 2017). It is unclear how the absence of the full functional complement of SA factors would affect SA vaccines in mice.

Conclusions and outlook

In contemplating the remarkable efficacy of the SARS-CoV-2 vaccines, one might almost be forgiven to think that developing vaccines is easy, and that the tried-and-true vaccinology toolbox still holds the key to successful SA vaccines. Yet vaccines have remained ineffective in repeated trials. Pathogenesis studies have informed us in so many ways that SA is different as a master of immune evasion strategies, and that induction of non-protective immunity is largely the rule. As suggested by leaders of the SA field, a more than incremental approach is now needed to address the fundamental root of vaccine failures. Bridging the translational divide will require simulation of the human host experience in

animal models. For example, SA vaccines could be studied in mice that have been infected or colonized previously with SA. Likewise, human anti-SA monoclonal antibodies could be tested in laboratory animals or humanized mice that have been infused with anti-SA antibodies purified from human sera. Our prediction is that, in both cases, vaccine efficacy would be significantly dampened in the modified experimental settings. If our hypothesis is validated, the findings would pave the way for more direct studies of the role of SA virulence determinants in vaccine modulation. Admittedly, no published studies to date have demonstrated a causal relationship between SA prior exposure and vaccine failures. We propose that these studies are urgently needed since understanding of SA reprogramming of the host environment and its effect on vaccination likely holds the clues to development of the next successful SA vaccine.

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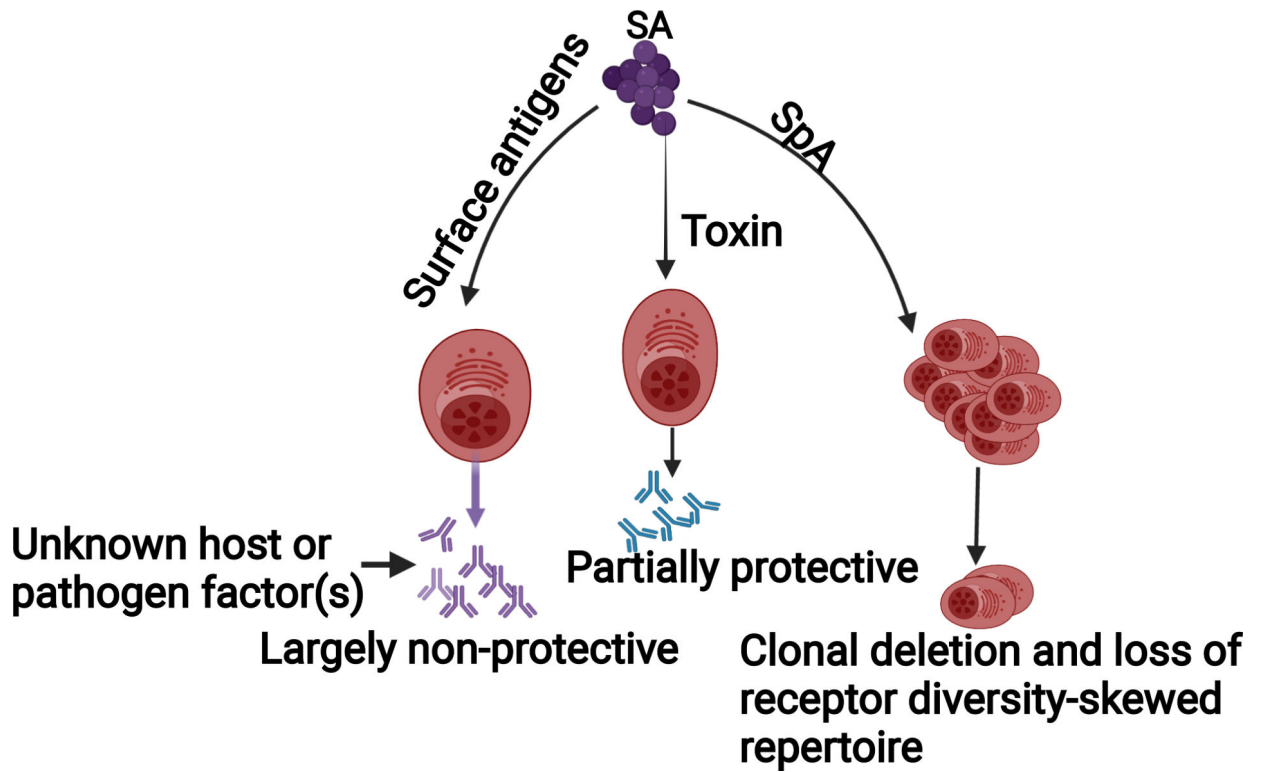


Figure 1. B cell evasion mechanisms.

Antibody responses against immunodominant SA surface antigens are largely non-protective, whereas responses against toxins are partially protective. SpA induces B cell deletion and suppression through its super-antigenic activity, and thus creates “holes” in the B cell repertoire and primes for a skewed specific antibody response. The reason why antibodies against surface antigens are not protective is unclear.

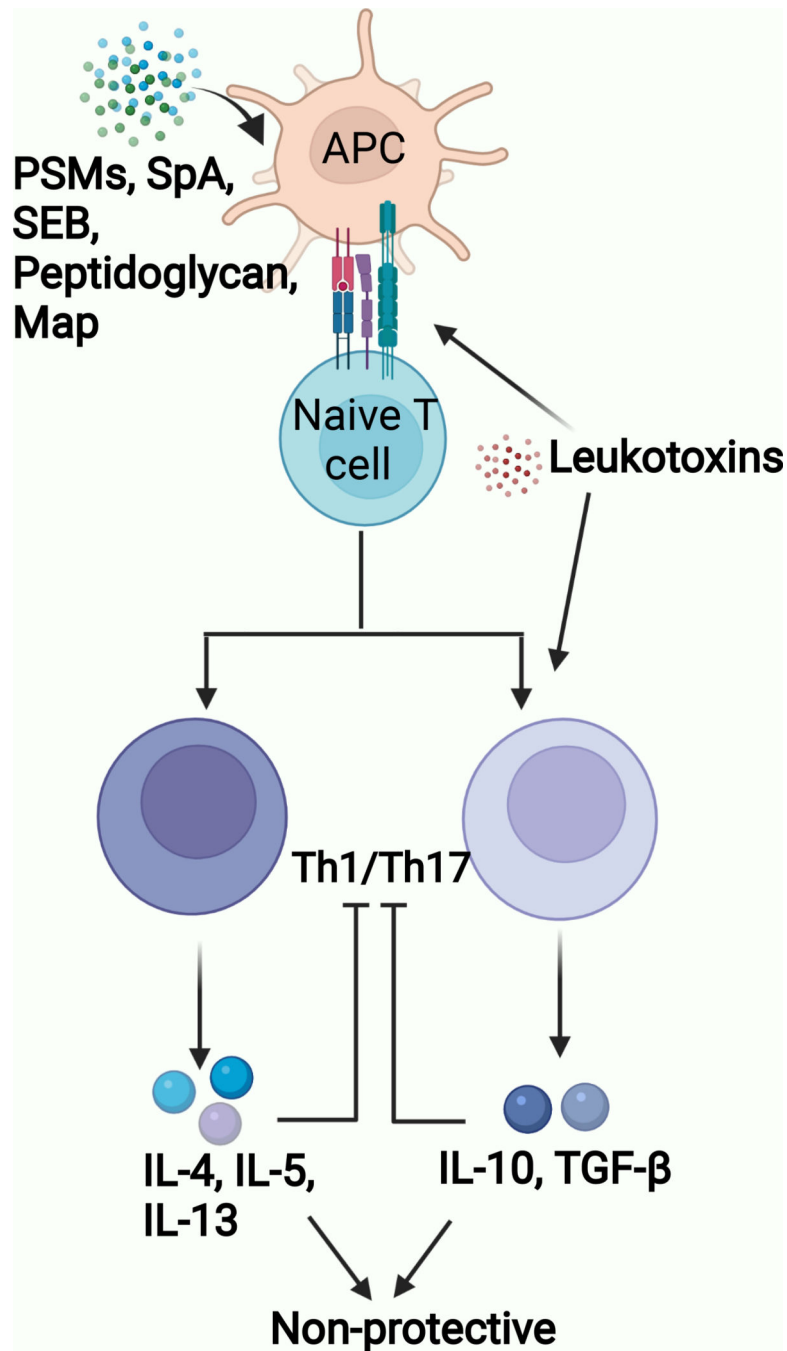


Figure 2. T cell evasion mechanisms.

Th1 / Th17 lymphocytes mediate *anti-SA* immunity. Staphylococcal toxins (α -toxin and LukED) induce cytolysis of mature and memory T cells. Various SA virulence determinants affect T cell priming by modulating antigen-presenting cell – naïve T cell interaction: PSM (induction of tolerogenic DC); SEB ($V\beta$ -specific T cell activation), Peptidoglycan modification (suppression of Th17-related cytokines), Map (induction of Th2 cells), SpA

(induction of Treg cells), LukED and LukAB (killing of DC). APC: Antigen Presenting Cells.

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