

Correlation between *In Vitro* Complement Deposition and Passive Mouse Protection of Anti-Pneumococcal Surface Protein A Monoclonal Antibodies

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The shortcomings of the licensed polysaccharide-based pneumococcal vaccine are driving efforts toward development of a protein-based vaccine that is serotype independent and effective in all age groups. An opsonophagocytic killing assay (OPKA) is used to evaluate the antibody response against polysaccharide-based pneumococcal vaccines. However, the OPKA is not reliable for noncapsular antigens. Thus, there is a need to develop an *in vitro* surrogate for protection for protein vaccine candidates like pneumococcal surface antigen A (PspA). PspA is a serologically variable cell surface virulence factor. Based on its sequence, PspA has been classified into families 1 (clade 1 and 2), 2 (clades 3, 4 and 5), and 3 (clade 6). Here, we report the characterization of 18 IgG anti-PspA monoclonal antibodies (anti-PspA^{hkR36A} MABs) generated from mice immunized with heat-killed strain R36A (clade 2). An enzyme-linked immunosorbent assay (ELISA)-based analysis of the reactivity of the MABs with recombinant PspAs from the 6 clades indicated that they were family 1 specific. This was confirmed by flow cytometry using a hyperimmune serum generated against PspA from R36A. Eight MABs that bind at least one clade 1- and clade 2-expressing strain were evaluated for complement deposition, bactericidal activity, and passive protection. The anti-PspA^{hkR36A} MAB-dependent deposition of complement on pneumococci showed a positive correlation with passive protection against strain WU2 ($r = 0.8783$, $P = 0.0041$). All of our protective MABs showed bactericidal activity; however, not all MABs that exhibited bactericidal activity conferred protection *in vivo*. The protective MABs described here can be used to identify conserved protection eliciting B cell epitopes for engineering a superior PspA-based vaccine.

The bacterial pathogen *Streptococcus pneumoniae* (pneumococcus) is responsible for causing pneumonia, septicemia, meningitis, and otitis media in humans (1). According to the estimate made by the World Health Organization in 2005, 1.6 million individuals die of diseases caused by *S. pneumoniae* every year, and most of these deaths occur in developing countries (2). In the year 2000, it was estimated that pneumococcal disease was responsible for about 800,000 deaths of children <5 years of age (3). The currently available pneumococcal polysaccharide vaccine is not effective in children <2 years of age. Pneumococcal conjugate vaccines, however, overcome this limitation and are effective in children but have limited serotype coverage (4). The development of antibiotic resistance and the emergence of nonvaccine serotypes pose difficulties in the management of pneumococcal infections. Efforts are being made globally to develop a protein-based pneumococcal vaccine that confers serotype-independent protection in all age groups (5–7).

A polysaccharide capsule envelops *S. pneumoniae*, and it serves as the major virulence factor by shielding pneumococci from immune attack. The unencapsulated strains of *S. pneumoniae* are known to be avirulent or highly attenuated. In addition to the capsule, several surface-associated proteins have been demonstrated to be involved in pneumococcal virulence, and one such protein is pneumococcal surface protein A (PspA) (8). A well-established opsonophagocytic killing assay (OPKA) is available for evaluating the pneumococcal polysaccharide-based vaccines. The recent interest in the protein-based pneumococcal vaccines has led to efforts toward development of an *in vitro* assay for noncapsular antigens that can help in predicting and quantitating the protective activity of antibodies against protein vaccine candidates. Various investigators have tried to correlate anti-protein

antibody titers, surface binding (by a whole-cell enzyme-linked immunosorbent assay [ELISA]), and a surface killing assay with *in vivo* protection (9–12). The notion of *in vitro* antibody-mediated complement deposition as a possible surrogate for predicting *in vivo* protection has been proposed by Goulart et al. and Ochs et al. (13, 14). However, these investigators did not validate it with *in vivo* protection experiments. Availability of a robust *in vitro* assay would help in minimizing the use of animal models for testing protein vaccine candidates.

PspA is a polymorphic, surface-associated choline-binding protein (15). PspA has a predominantly α -helical coiled coil structure (16, 17). It is present in essentially all clinical isolates studied to date and is being pursued as a promising vaccine candidate (18). Based on the amino acid sequence, PspA has been classified

Received 28 December 2013 Returned for modification 3 February 2014

Accepted 11 November 2014

Accepted manuscript posted online 19 November 2014

Citation Khan N, Qadri RA, Sehgal D. 2015. Correlation between *in vitro* complement deposition and passive mouse protection of anti-pneumococcal surface protein A monoclonal antibodies. *Clin Vaccine Immunol* 22:99–107. doi:10.1128/CVI.00001-14.

Editor: T. S. Alexander

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Supplemental material for this article may be found at <http://dx.doi.org/10.1128/CVI.00001-14>.

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doi:10.1128/CVI.00001-14

into three families and six clades. Family 1 includes clades 1 and 2, family 2 includes clades 3, 4, and 5, and family 3 includes clade 6 (19). Studies have shown that 94 to 99% of the pneumococcal isolates analyzed belong to PspA families 1 and 2 (20, 21).

The complement-mediated clearance of pneumococci is an important component of the host defense mechanism (22). A PspA-deficient strain is cleared faster than wild-type pneumococci, and an anti-PspA antibody facilitates complement-dependent phagocytosis of *S. pneumoniae* (23). Ren and coworkers have demonstrated that anti-PspA antibodies enhance complement activation and deposition on pneumococcal surface and thus help in clearance (24).

Active immunization with PspA in animal models has proven to be protective against invasive disease and nasopharyngeal carriage (25). Mice immunized with DNA vaccine expressing the extracellular domain of PspA were protected against an intraperitoneal challenge with a pneumococcal strain bearing PspA from the same clade (26). The B cell epitopes recognized by protective monoclonal antibodies (MAbs) have been mapped to the 192- to 260-amino acid region (27). Daniels et al. recently demonstrated that the proline-rich region of PspA contains surface-accessible epitopes that are protective in both active and passive mouse protection experiments (28). PspA has been shown to elicit high antibody titers in humans, and human anti-PspA sera can protect mice against pneumococcal challenge when transferred passively (18, 29).

There is evidence to suggest that not all anti-PspA antibodies are protective. The goals of the present study were to identify anti-PspA^{hkr36A} MAbs that recognize conserved cross-protective B cell epitopes, and since the *in vitro* surrogate of protection is not well established for noncapsular antigens, we evaluated the surface binding, complement deposition, and bactericidal activity of anti-PspA^{hkr36A} MAbs as potential *in vitro* correlates of protection. We found that all of the 18 anti-PspA^{hkr36A} MAbs recognized family 1 PspAs and did not bind PspAs representing families 2 and 3. We identified 4 anti-PspA^{hkr36A} MAbs (P1E11, M4F4, P2A4, and P2B5) that augmented complement deposition on pneumococci, exhibited bactericidal activity, and conferred protection against PspA clade 1- and 2-bearing *S. pneumoniae* strains in passive mouse protection experiments. Further, our data with strain WU2 suggested that anti-PspA^{hkr36A} MAb-dependent complement deposition on pneumococci strongly correlated with *in vivo* protection. We observed that all of the protective anti-PspA^{hkr36A} MAbs exhibited bactericidal activity; however, not all of the anti-PspA^{hkr36A} MAbs that showed bactericidal activity conferred *in vivo* protection.

MATERIALS AND METHODS

Mice. Six- to 8-week-old BALB/c (female) and CBA/N (male/female) inbred strains of mice were obtained from the Small Animal Facility of the National Institute of Immunology. Animals were rested and handled in accordance with the institutional animal ethics committee guidelines. Blood samples from healthy donors were taken with the approval of and following the guidelines of the institutional human ethics committee. Experiments involving recombinant DNA and handling of *S. pneumoniae* were carried out in accordance with the institutional biosafety committee guidelines.

Pneumococcal strains, plasmids, and culture conditions. The pneumococcal strains and plasmids used in this study are listed in Table S1 in the supplemental material. Pneumococcal strains were maintained in Todd-Hewitt broth supplemented with 0.5% yeast extract (THY) or on a

plate with tryptic soy agar (TSA) supplemented with 5% (vol/vol) sheep blood at 37°C in the presence of 5% CO₂. The mid-logarithmic-phase pneumococcal cultures were stored with 15% (vol/vol) glycerol or 17% fetal calf serum, aliquoted, and stored at -70°C as described previously (30).

Escherichia coli cells were maintained in Luria-Bertani broth or on a Luria-Bertani agar plate with antibiotic(s) wherever required.

Molecular cloning, overexpression, and purification of recombinant PspA. The subfragments encoding the N-terminal (surface exposed) region of PspA for clade 3 (PspA^{TIGR4}) and clade 5 (PspA^{ATCC 6303}) were amplified by PCR using genomic DNA from the pneumococcal strains TIGR4 and ATCC 6303, respectively, following the cloning strategy and PCR conditions described by Rohatgi et al. (30). The corresponding plasmid constructs for PspA clade 1 (pUAB069, strain L82016 [PspA^{L82016}]), clade 4 (pUAB100, strain JCP#56 [PspA^{JCP#56}]), and clade 6 (pUAB104, strain BG9300 [PspA^{BG9300}]) were kindly provided by Susan Hollingshead, University of Alabama, USA. The plasmid construct for PspA clade 2 (R36A [PspA^{R36A}]) was published previously from our laboratory (30). For expression purposes, pQE-30 Xa- and pET-20b-based constructs were transformed into *E. coli* expression strains SG13009 and BL-21(DE3), respectively. Recombinant PspA was purified using nickel-nitrilotriacetic acid (Ni-NTA) affinity chromatography (Sigma-Aldrich, USA). The purity of the protein preparation was found to be >95% by SDS-PAGE and was of the expected molecular size.

Enzyme-linked immunosorbent assay. The reactivities of the 18 IgG anti-PspA^{hkr36A} MAbs with recombinant PspAs representing the six clades of PspA were analyzed by an ELISA. Briefly, 96-well polystyrene microtiter plates (Greiner Bio-One, Germany) were coated overnight at 4°C with recombinant PspA (50 µl of 2 µg/ml per well) in 100 mM carbonate-bicarbonate buffer (pH 9.5). The plates were washed with phosphate-buffered saline (PBS) containing 0.05% Tween 20 (PBST) and blocked with PBS containing 2% bovine serum albumin (BSA) at 37°C for 1 h. After washing with PBST, the plates were incubated with the culture supernatant from the 18 anti-PspA^{hkr36A} hybridomas (in duplicate) at 37°C for 1 h. The plates were washed with PBST and incubated with horseradish peroxidase-conjugated goat anti-mouse Ig antibody (diluted 1 in 2,500; Becton-Dickinson Bioscience, USA) followed by incubation at 37°C for 1 h. The color was developed using 3,3',5,5'-tetramethylbenzidine-H₂O₂ as the substrate, and absorbance was recorded at 450 nm.

Generation of polyclonal sera against recombinant PspA^{R36A}. Six- to 8-week-old female BALB/c mice were immunized subcutaneously with 25 µg recombinant PspA^{R36A} emulsified with Imject alum (1:1 [wt/wt]) (Pierce, USA). On days 14 and 28, mice were given a booster injection with the same amount of antigen emulsified as described above. The control mice received only Imject alum in PBS. One week after the second booster, hyperimmune serum was isolated after mice were bled retro-orbitally, and an ELISA was performed to determine the PspA-specific antibody titer.

Surface staining with anti-PspA^{hkr36A} MAbs and anti-PspA^{R36A} hyperimmune serum. The surface binding of anti-PspA^{hkr36A} MAbs with *S. pneumoniae* strains that express clade 1 PspA (BG8838) and clade 2 PspA (WU2 and D39) was analyzed using flow cytometry as described previously (31). Briefly, mid-logarithmic-phase (optical density at 600 nm [OD₆₀₀] of 0.4) pneumococci (10⁷ CFU) were washed with PBS and incubated with 200 µl of culture supernatant from the hybridomas at room temperature for 1 h. After washing with PBS, pneumococci were incubated with fluorescein isothiocyanate (FITC)-conjugated F(ab')₂ fragment goat anti-mouse IgG plus IgM(H+L) antibody (diluted 1 in 200) (Jackson ImmunoResearch Laboratory, USA) followed by incubation at room temperature for 1 h. After washing, pneumococci were fixed with 2% paraformaldehyde (PFA) for 10 min at 4°C, and surface staining was analyzed by flow cytometry (FACSCalibur; Becton-Dickinson Bioscience).

The surface binding of the anti-PspA^{R36A} polyclonal sera (diluted 1 in 200) was analyzed with strains BG8838 (clade 1), D39 (clade 2), TIGR4 (clade 3), EF5668 (clade 4), ATCC 6303 (clade 5), and BG6803 (clade 6)

TABLE 1 ELISA-based analysis for reactivity of anti-PspA^{hkr36A} MAbs with recombinant PspAs representing the 6 PspA clades^a

Hybridoma	Fold change for pneumococcal strain (PspA family/clade):					
	L82016 (1/1) ^b	R36A (1/2) ^b	TIGR4 (2/3)	JCP#56 (2/4)	ATCC 6303 (2/5)	BG9300 (3/6)
B3D12	16.5	17.2	0.8	0.8	0.8	1.2
B3H8	1.8	2.1	1.0	0.9	1.0	0.8
L5C8	17.1	16.7	0.8	0.8	0.8	0.8
L5F10	19.5	26.9	0.8	0.7	0.9	1.2
M4F4	30.0	32.5	1.1	1.2	1.6	1.2
P1E11	29.2	31.5	1.4	1.4	1.7	1.6
D1A5	10.6	11.9	0.9	0.8	0.7	0.9
K1B12	15.9	16.5	0.7	0.7	0.6	0.9
M6B2	13.5	14.5	1.0	1.4	1.3	1.9
P2F9	9.8	13.4	0.9	1.0	0.9	1.0
P2A4	21.6	19.9	1.0	1.5	0.9	1.0
P2B5	22.0	24.0	1.0	1.2	1.2	1.2
J4C1	28.2	29.3	1.0	0.8	0.7	0.8
P2C2	16.2	17.0	0.6	0.6	0.6	0.6
A1D9	2.0	2.7	0.6	0.7	0.7	0.6
C4B4	18.1	18.1	0.8	0.6	0.6	0.7
F4B6	24.7	26.3	1.0	0.9	0.8	1.1
D3H6	4.0	5.7	0.7	0.7	0.7	1.1

^a The culture supernatants from the 18 anti-PspA^{hkr36A} MAb-secreting hybridomas were tested for reactivity with recombinant PspAs representing the 6 clades of PspA (extracellular domain with or without the proline-rich region) by an ELISA. The numerical values represent fold change in absorbance relative to the control.

^b These columns show fold change values that were greater than twice the value obtained with the control and were considered significant.

using flow cytometry as described above. Geometric mean fluorescence intensity (GMFI) values equal to or greater than twice the value obtained with the preimmune control were considered significant.

Purification of anti-PspA^{hkr36A} MAbs. Hybridomas secreting anti-PspA^{hkr36A} MAbs were generated and cultured as previously described (30). Ascitic fluid was generated for the anti-PspA^{hkr36A} MAb-secreting hybridomas. Briefly, BALB/c mice (3 mice per hybridoma) were injected intraperitoneally with 0.5 ml incomplete Freund's adjuvant (Sigma-Aldrich). Three days later, 5 × 10⁶ hybridoma cells were injected intraperitoneally. After 7 to 10 days, ascitic fluid from the peritoneal cavity was centrifuged at 500 × g for 10 min at 4°C, and the supernatant was collected and stored at -70°C. The MAbs were purified from ascitic fluid using protein G-Sepharose beads (GE Healthcare, USA) as described earlier (32). The protein concentration of the purified MAb was estimated using a micro BCA protein assay kit (Thermo-Scientific, USA).

Passive mouse protection assay. Groups of eight 6- to 8-week-old CBA/N mice were injected intraperitoneally with purified anti-PspA^{hkr36A} MAb or a matched isotype control. The amounts of MAb administered for the high- and low-dose experiments were 5 and 1.25 mg/kg body weight, respectively. An hour later, mice were challenged intravenously with 10⁷ CFU (100 times the 50% lethal dose [LD₅₀]) of strain BG8838 and 10³ CFU (100 times the LD₅₀) of WU2 (27). The survival of mice was monitored every 12 h for the first 10 days and every 24 h for the next 11 days.

Blood bactericidal assay. The anti-PspA^{hkr36A} MAbs were analyzed by a blood bactericidal assay as described previously (33). Briefly, human peripheral blood was collected using recombinant hirudin from yeast (100 U/ml blood) as an anticoagulant. Pneumococci (500 CFU in 10 μl) were incubated with 235 μl of blood in the presence of either purified anti-PspA^{hkr36A} MAb or the corresponding isotype control (5 μl of 1 mg/ml). Samples were incubated at 37°C with rotation for 2 h for D39 and 3 h for BG8838. The surviving bacteria were enumerated by plating serial dilutions (in duplicate) on TSA plates, and the mean values obtained were used to calculate bactericidal activity as described below. The data are

TABLE 2 Surface binding of anti-PspA^{hkr36A} MAbs with strains expressing family 1 PspA^a

MAb	GMFI (fold change) for:		
	BG8838 clade 1	WU2 clade 2	D39 clade 2
IgG1 IC	4.77 (1.0)	3.75 (1.0)	3.67 (1.0)
B3D12	13.10 (2.8)	5.18 (1.4)	64.80 (17.7)
B3H8	14.70 (3.1)	5.49 (1.5)	8.75 (2.4)
L5C8	18.70 (3.9)	3.82 (1.0)	137.00 (37.3)
L5F10	18.90 (4.0)	7.22 (1.9)	66.20 (18.1)
M4F4	59.50 (12.5)	11.80 (3.2)	140.00 (38.2)
P1E11	81.80 (17.2)	14.20 (3.8)	146.00 (39.8)
D1A5	4.04 (0.8)	107.00 (28.5)	52.90 (14.4)
K1B12	3.86 (0.8)	9.39 (2.5)	69.80 (19.0)
M6B2	5.64 (1.2)	19.20 (5.1)	174.00 (47.4)
P2F9	4.80 (1.0)	4.12 (1.1)	14.30 (3.9)
IgG2a IC	4.82 (1.0)	2.82 (1.0)	4.68 (1.0)
P2A4	23.60 (4.9)	11.10 (3.9)	59.70 (12.8)
P2B5	17.60 (3.7)	16.10 (5.7)	33.90 (7.3)
J4C1	3.88 (0.8)	3.27 (1.2)	60.80 (13.0)
P2C2	3.52 (0.7)	8.93 (3.2)	34.80 (7.4)
IgG2b IC	4.16 (1.0)	2.28 (1.0)	3.21 (1.0)
A1D9	4.35 (1.1)	16.3 (7.2)	47.40 (14.8)
C4B4	10.80 (2.6)	22.10 (9.7)	45.00 (14.1)
F4B6	3.70 (0.89)	56.7 (24.9)	41.00 (12.8)
IgG3 IC	4.26 (1.0)	2.82 (1.0)	3.21 (1.0)
D3H6	4.41 (1.0)	4.32 (1.5)	25.80 (8.1)

^a Surface binding of 18 anti-PspA^{hkr36A} MAbs with strains BG8838 (clade 1) and WU2 and D39 (clade 2) was analyzed by flow cytometry. Strain D39 was included in the analysis as the anti-PspA MAbs were raised against R36A, an unencapsulated derivative of D39. Pneumococci were incubated with 200 μl of culture supernatant from the 18 anti-PspA^{hkr36A} MAb-secreting hybridomas followed by staining with the appropriate FITC-conjugated secondary antibody. Matched isotype control MAbs were included, and samples were analyzed using a flow cytometer. The surface binding is expressed as geometric mean fluorescence intensity (GMFI), and the fold change relative to the corresponding isotype control MAb (assigned 1.0) is given in parentheses. A ≥2-fold increase in the surface staining relative to that of the corresponding isotype control was considered significant. IgG1 IC, IgG2a IC, IgG2b IC, and IgG3 IC represent isotype control MAbs for the isotypes IgG1, IgG2a, IgG2b, and IgG3, respectively.

presented as percent killing, which was arrived at using the formula ([colony count with appropriate isotype control MAb - colony count with anti-PspA^{hkr36A} MAb] divided by colony count with the appropriate isotype control MAb) × 100.

Complement deposition assay. The complement deposition assay was performed as described by Ochs et al. with some modifications (14). Briefly, mid-logarithmic-phase pneumococci (10⁷ CFU) were washed with PBS and incubated with either an anti-PspA^{hkr36A} MAb or a matched isotype control (100 μl of 20 μg/ml) for 1 h at room temperature. Treated pneumococci were incubated in 200 μl of normal human serum pooled from three healthy donors (diluted to 10% in Hanks' balanced salt solution with Ca²⁺ and Mg²⁺ ions) (Biological Industries, Israel) for 30 min at 37°C. The mixture was incubated on ice for 10 min and washed with PBS, and the incubation was continued with PBS-1% BSA containing mouse anti-human C3 MAbs (diluted 1 in 50) (Abcam, USA) for 45 min on ice. The bound anti-C3 antibody was detected using an FITC-conjugate F(ab')₂ fragment goat anti-mouse IgG plus IgM(H+L) antibody (diluted 1 in 200). Pneumococci were fixed with 2% PFA for 10 min on ice and analyzed by flow cytometry.

Statistical analysis. Statistical analysis was done using GraphPad Prism version 6 software (GraphPad Software Inc., USA). The passive mouse protection data were compared using the log rank test. We used a

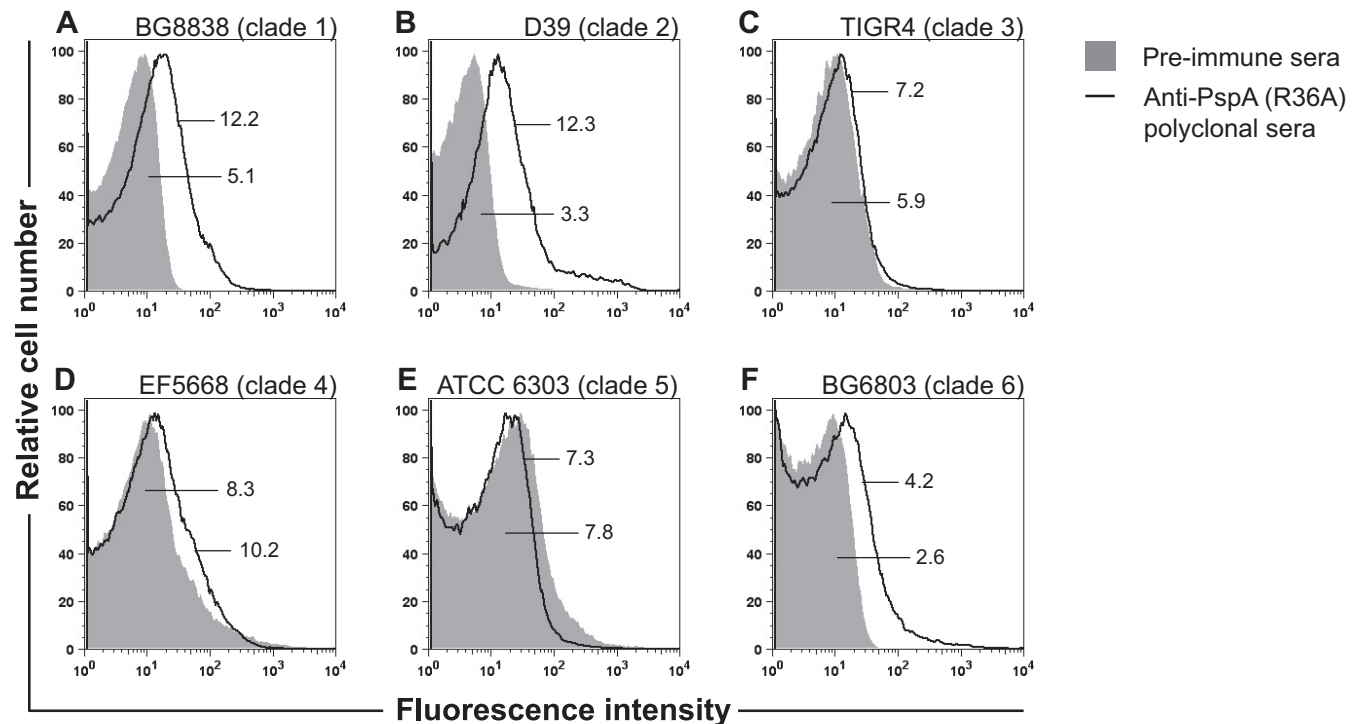


FIG 1 Surface binding of anti-PspA^{R36A} polyclonal sera with pneumococcal strains representing the 6 clades of PspA. Surface binding with strains BG8838 (A), D39 (B), TIGR4 (C), EF5668 (D), ATCC 6303 (E), and BG6803 (F) was analyzed by flow cytometry using anti-PspA^{R36A} polyclonal and preimmune sera. Preimmune serum (solid gray histogram) was used as the negative control. The GMFI values for the anti-PspA^{R36A} polyclonal and preimmune sera are shown. GMFI values of ≥ 2 times the value obtained with the preimmune sera were considered significant.

parametric Pearson's linear regression correlation analysis to evaluate a possible correlation between the anti-PspA^{hkr36A} MAb-dependent deposition of complement on pneumococci and passive mouse protection. A *P* value of <0.05 was considered statistically significant.

RESULTS

Cross-reactivity and surface staining of anti-PspA^{hkr36A} MAbs is restricted to PspA family 1. We previously reported that the relative avidity of the primary IgG anti-PspA polyclonal antibody was higher than that of recombinant PspA^{R36A} in the sera of mice immunized with heat-killed R36A (30). In this study, we analyzed 18 IgG anti-PspA^{hkr36A} MAbs for the extent of cross-reactivity with recombinant PspA belonging to the 6 PspA clades (i.e., PspA^{L82016}, PspA^{R36A}, PspA^{TIGR4}, PspA^{ICP#56}, PspA^{ATCC 6303}, and PspA^{BG9300}) by an ELISA (Table 1). All of the 18 MAbs reacted with PspAs representing clades 1 and 2 (family 1). None of the MAbs bound PspAs representing families 2 (clades 3, 4, and 5) and 3 (clade 6), suggesting that the anti-PspA^{hkr36A} MAbs were family 1 specific.

Next, we assessed whether these 18 anti-PspA^{hkr36A} MAbs bind to the surface of pneumococci. For this purpose, we tested strains BG8838 (clade 1) and WU2 and D39 (clade 2). Of the 18 MAbs, BG8838 and WU2 were recognized by 9 and 10 MAbs, respectively (Table 2). As expected, all 18 anti-PspA MAbs stained D39. Nine MAbs bound one clade 1 and two clade 2 PspA-expressing strains. Eight (i.e., B3D12, B3H8, L5C8, L5F10, M4F4, P1E11, P2A4, and P2B5) of these were selected at random for further analysis.

To confirm our observation that the anti-PspA antibody response elicited against heat-killed R36A was family 1 specific, the anti-PspA^{R36A} hyperimmune serum was used to test surface bind-

ing with strains expressing PspAs belonging to families 1, 2, and 3 by flow cytometry (Fig. 1). The data demonstrate that anti-PspA^{R36A} hyperimmune serum binds to family 1 PspA-expressing strains BG8838 (clade 1) and D39 (clade 2). No binding was observed with PspA family 2-bearing strains TIGR4 (clade 3), EF5668 (clade 4), and ATCC 6303 (clade 5) (Fig. 1). Similarly, there was no surface staining with the PspA family 3-expressing strain BG6803 (clade 6). Thus, our flow cytometry data demonstrate that the binding of anti-PspA^{R36A} hyperimmune serum to pneumococcal strains was family 1 specific, and this corroborates the data obtained with anti-PspA^{hkr36A} MAbs (Table 2).

In vivo protective efficacy of anti-PspA^{hkr36A} MAbs. The relative protective activities of anti-PspA^{hkr36A} MAbs B3D12, B3H8, L5C8, L5F10, M4F4, P1E11, P2A4, and P2B5 were assessed by a passive mouse protection assay using BG8838 (clade 1) and WU2 (clade 2) as the challenge strains. CBA/N mice were given either purified anti-PspA^{hkr36A} or the corresponding isotype control MAb (high dose; 5 mg/kg body weight) intraperitoneally. One hour later, mice were challenged intravenously with either purified anti-PspA^{hkr36A} or the corresponding isotype control MAb (high dose; 5 mg/kg body weight) intraperitoneally. One hour later, mice were challenged intravenously with either BG8838 or WU2. The mouse survival data with BG8838 as the challenge strain (Fig. 2A, C, and E) suggested that anti-PspA^{hkr36A} MAbs M4F4, P1E11, and L5C8 provided 87.5, 75, and 50% protection, respectively, compared to that for the corresponding isotype control (12.5%) (Fig. 2A). The MAbs L5F10, B3H8, and B3D12 provided 75, 62.5 and 37.5% protection, respectively, when challenged with BG8838 (Fig. 2C). All mice in the set that received the IgG1 isotype control MAb died within 7 days. The MAbs P2A4 and P2B5 protected 100 and 62.5% of the mice, respectively, when challenged with BG8838, compared to 12.5% in

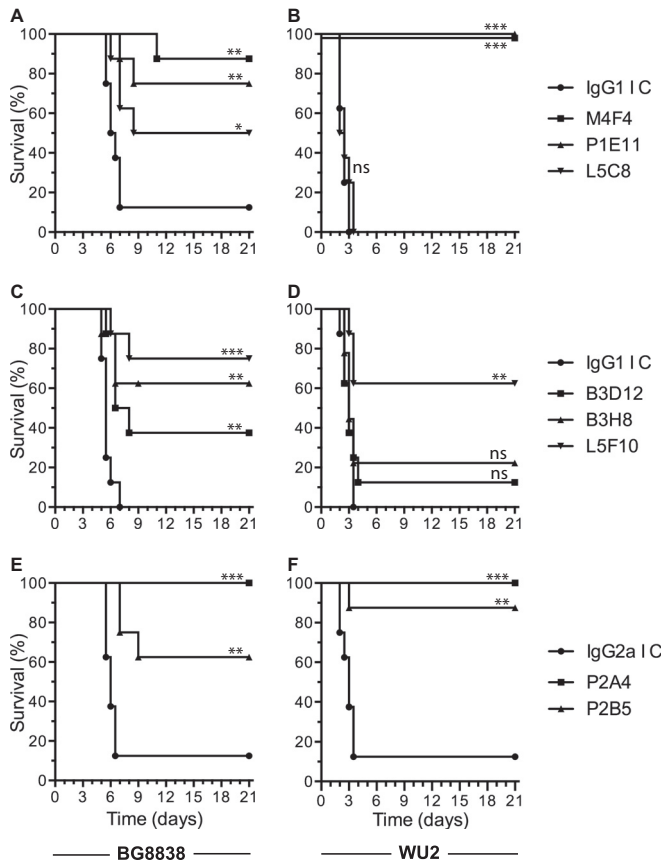


FIG 2 Relative efficacy of anti-PspA^{hkR36A} MAbs to protect mice against intravenous challenge. CBA/N mice were injected with purified anti-PspA^{hkR36A} MAb M4F4, P1E11, or L5C8 (A and B), B3D12, B3H8, or L5F10 (C and D) or P2A4 or P2B5 (E and F) intraperitoneally at 5 mg/kg body weight (high dose). The corresponding isotype control MAb (IgG1 IC or IgG2a IC) was included in each set as the negative control. One hour later, mice were challenged with 10⁷ CFU of BG8838 (A, C, and E) or 10³ CFU of WU2 (B, D, and F), and mouse survival was recorded. The data for the group given anti-PspA^{hkR36A} MAb were compared with those for the respective isotype control MAb using the log rank test. *, P < 0.05; **, P < 0.01; ***, P < 0.001; ns, statistically not significant.

the case of the isotype control (Fig. 2E). In experiments where mice were challenged with WU2 (Fig. 2B, D, and F), M4F4 and P1E11 provided 100% protection. L5C8 failed to confer any protection, like the IgG1 isotype control MAb (Fig. 2B). L5F10 conferred 62.5% protection and B3H8 and B3D12 provided ≤25% protection when mice were challenged with WU2 (Fig. 2D). All of the mice in the corresponding control group died within 3.5 days. P2A4 provided 100% and P2B5 provided 87.5% protection when mice were challenged with WU2, whereas the corresponding IgG2a isotype control MAb protected only 12.5% of the mice (Fig. 2F).

On the basis of results of the passive mouse protection experiments described above, MAbs M4F4, P1E11, L5F10, P2A4, and P2B5 were selected as they conferred >50% protection against the 2 challenge strains tested. We repeated the experiments with a lower dose (1.25 mg/kg body weight) of the purified anti-PspA^{hkR36A} MAbs and the corresponding isotype control. The mouse survival data following challenge with BG8838 suggested that P1E11, M4F4, and L5F10 provided 75, 62.5, and 37.5% protection, respectively, whereas all mice in the control group died

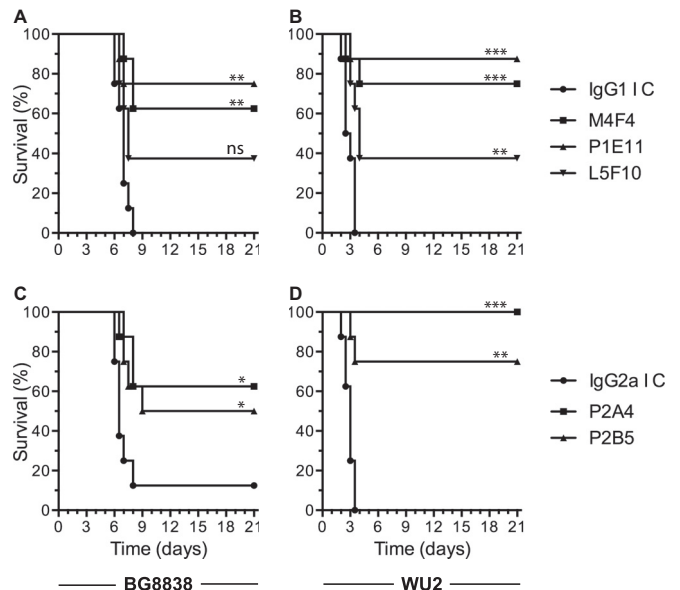


FIG 3 Anti-PspA^{hkR36A} MAbs protect mice against pneumococcal infection even when given at a lower dose. CBA/N mice were injected intraperitoneally with 1.25 mg/kg body weight (low dose) of either anti-PspA^{hkR36A} MAb M4F4, P1E11, or L5F10 (A and B) or P2A4 or P2B5 (C and D). The control group was given the respective isotype control MAb (IgG1 IC or IgG2a IC). Mice were challenged with BG8838 (A and C) or WU2 (B and D) 1 h later, and mouse survival was recorded. For other details, refer to the legend to Fig. 2.

within 8 days (Fig. 3A), while P2A4 and P2B5 provided 62.5 and 50% protection, respectively, compared to the IgG2a isotype control for which 12.5% survival was observed (Fig. 3C). The mouse survival data with WU2 as the challenge strain suggested that P1E11, M4F4, and L5F10 provided 87.5, 75, and 37.5% protection, respectively (Fig. 3B), while P2A4 and P2B5 provided 100 and 75% protection, respectively (Fig. 3D). In both of the experiments, all control mice died within 3.5 days. The consolidated data from the passive protection experiments demonstrate that 4 anti-PspA^{hkR36A} MAbs (M4F4, P1E11, P2A4, and P2B5) confer ≥50% protection against intravenous challenge with PspA clade 1- and 2-expressing strains.

Correlation between *in vivo* protection and *in vitro* complement deposition and bactericidal activity. We were interested in determining the possible correlation between *in vivo* protection and *in vitro* bactericidal activity and the complement deposition capability of anti-PspA^{hkR36A} MAbs. We employed a blood bactericidal assay to evaluate the ability of the 8 anti-PspA^{hkR36A} MAbs (i.e., B3D12, B3H8, L5C8, L5F10, M4F4, P1E11, P2A4, and P2B5) to kill the PspA family 1-expressing strains BG8838 (clade 1) and D39 (clade 2). Pneumococci were incubated with blood from healthy donors in the presence of either purified anti-PspA^{hkR36A} or a matched isotype control MAb, and the surviving pneumococci were enumerated by plating. The results suggest that all 8 anti-PspA^{hkR36A} MAbs exhibited significant bactericidal activity (ranging from 25.7 to 80.3% compared with that of the matched isotype control) against BG8838 and D39, although the extent of bactericidal activity varied from one MAb to the other (Fig. 4).

We next tested whether these MAbs can enhance C3 deposition on the surface of clade 1 (BG8838)-expressing and clade 2 (WU2 and D39)-expressing strains (Table 3). Monoclonal antibodies

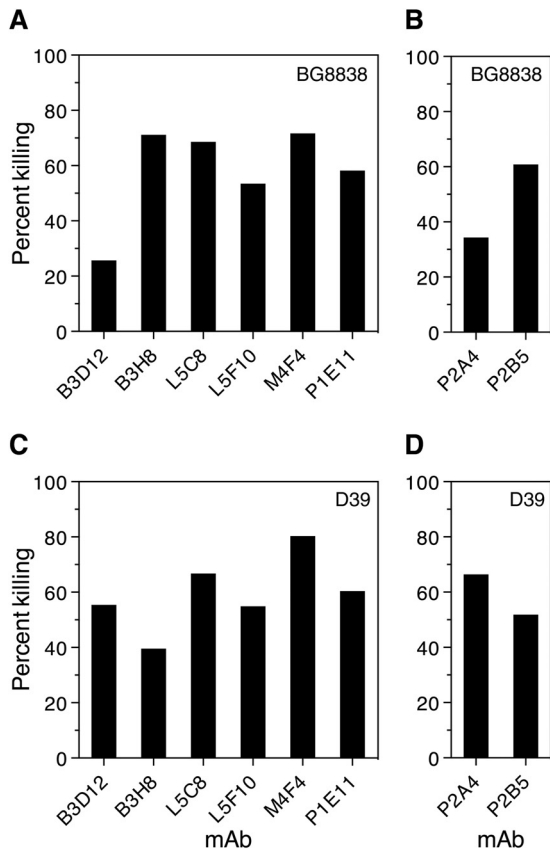


FIG 4 Anti-PspA^{hkr36A} MAbs promote pneumococcal killing by human blood *in vitro*. *S. pneumoniae* strains BG8838 (A and B) and D39 (C and D) were incubated with blood from healthy donors and purified anti-PspA^{hkr36A} or a matched isotype control MAb. Matched IgG1 and IgG2a isotype control MAbs were used as the comparator in panels A and C and B and D, respectively (not shown). The contents were rotated for (3 h for BG8838 and 2 h for D39) at 37°C, and the surviving pneumococci were enumerated by plating. The identity of the anti-PspA^{hkr36A} MAb and the percent killing are plotted on the x and y axes, respectively. The percent bacterial killing was calculated as described in Materials and Methods. The assay was performed at least four times, and data from a single representative experiment are shown.

M4F4, P1E11, P2A4, and P2B5 showed significant enhancement in complement deposition on pneumococci compared to that of the corresponding isotype control for the 3 strains analyzed. Seven of the 8 MAbs augmented complement deposition on D39 with B3H8 being the exception. B3H8, however, enhanced deposition of complement on WU2. It was observed that the extent of the enhancement in C3 deposition depended on the anti-PspA^{hkr36A} MAb and the target strain.

We wanted to find out which aspect or feature of the antibody response correlated with *in vivo* protection. The surface staining, complement deposition, bactericidal activity, and mouse protection data for MAbs B3D12, B3H8, L5C8, L5F10, M4F4, P1E11, P2A4, and P2B5 are summarized in Table 4. The trend from the data appears to be that the higher the extent of complement deposition, the higher the bactericidal activity and *in vivo* protection. This is well illustrated by MAbs M4F4, P1E11, P2A4, and P2B5. While all the protective MAbs showed bactericidal activity, not all MAbs that exhibited bactericidal activity showed passive protection. For example, MAbs B3D12, B3H8, and L5C8 exhibited significant bactericidal activity but showed no to poor ability to confer protection when given passively to mice.

To determine whether *in vitro* MAb-dependent deposition of complement correlated with passive mouse protection, a parametric Pearson’s linear regression correlation analysis was performed. The analysis indicated that there was a positive correlation between *in vitro* complement deposition and the passive protection for strains WU2 (Fig. 5) and BG8838 (data not shown). The correlation was highly statistically significant for strain WU2 ($r = 0.8783$, $P = 0.0041$). Our data suggest that the antibody-dependent deposition of complement on the pneumococcal surface can be a potential *in vitro* correlate of protection for strains like WU2.

DISCUSSION

PspA is a likely candidate for a protein-based vaccine against pneumococcal infections; however, its serological variability might restrict the coverage of a PspA-based vaccine. For this reason, gaining insight into the nature of the variability of PspAs has been the subject of several studies that are directed at development of a protein-based pneumococcal vaccine. Studies aimed to investigate the level of cross-reactivity among PspAs in mice indicate that antibodies generated against PspA show higher cross-reactivity with the strains expressing PspA of the same family than the

TABLE 3 Flow cytometry-based analysis of complement C3 deposition on the surface of pneumococci in the presence of anti-PspA^{hkr36A} MAbs^a

Strains (clade)	GMFI (fold change) for:											
	Negative control ^b	Positive control ^c	IgG1 isotype control	B3D12 ^d	B3H8 ^d	L5C8 ^d	L5F10 ^d	M4F4 ^d	P1E11 ^d	IgG2a isotype control	P2A4 ^e	P2B5 ^e
BG8838 (1)	4.5	16.6	15.3	29.6	30.3	24.0	14.4	71.7 ^f	233.0 ^f	12.3	39.4 ^f	106.0 ^f
WU2 (2)	3.0	3.4	3.2	3.8	20.5 ^f	3.4	5.2	482.0 ^f	442.0 ^f	3.6	320.0 ^f	391.0 ^f
D39 (2)	5.5	7.1	7.0	418.0 ^f	12.9	516.0 ^f	511.0 ^f	568.0 ^f	523.0 ^f	6.1	429.0 ^f	384.0 ^f

^a The PspA family 1-expressing strains BG8838 (clade 1) and WU2 and D39 (clade 2) were incubated with either anti-PspA^{hkr36A} or a matched isotype control MAb. The bound complement C3 was detected by flow cytometry using anti-human C3 antibody followed by the appropriate FITC-conjugated secondary antibody.

^b Pneumococci were incubated with anti-human complement C3 antibody followed by the FITC-conjugated secondary antibody.

^c Pneumococci were incubated with 10% pooled normal human serum in Hanks’ balanced salt solution followed by anti-human C3 antibody and the FITC-conjugated secondary antibody.

^d These anti-PspA^{hkr36A} MAbs are of the IgG1 isotype.

^e These anti-PspA^{hkr36A} MAbs are of the IgG2a isotype.

^f GMFI values greater than or equal to twice the value obtained with the corresponding isotype control were considered significant.

TABLE 4 *In vivo* protective efficacies of anti-PspA^{hkR36A} MAbs correlate with the extent of complement deposition^a

Experiment	Strain	Result for anti-PspA ^{hkR36A} MAb:								
		B3D12	B3H8	L5C8	L5F10	M4F4 ^b	P1E11 ^b	P2A4 ^b	P2B5 ^b	
Surface staining (x) ^c	BG8838	2.8	3.1	3.9	4.0	12.5	17.2	4.9	3.7	
	WU2	1.4	1.5	1.0	1.9	3.2	3.8	3.9	5.7	
	D39	17.7	2.4	37.3	18.1	38.2	39.8	12.8	7.3	
Complement deposition (x) ^c	BG8838	1.9	2.0	1.6	0.9	4.7	15.2	3.2	8.6	
	WU2	1.2	6.4	1.1	1.6	150.6	138.1	88.9	108.6	
	D39	59.7	1.8	73.7	73.0	81.1	74.7	70.3	62.9	
Bactericidal activity (%) ^d	BG8838	25.7	71.2	68.6	53.5	71.7	58.2	34.3	60.8	
	D39	55.4	39.6	66.8	54.8	80.3	60.4	66.5	51.9	
Mouse passive protection assay (%) ^e	5 mg/kg	BG8838	37.5	62.5	50.0	75.0	87.5	75.0	100.0	62.5
		WU2	12.5	22.5	0.0	62.5	100.0	100.0	100.0	87.5
	1.25 mg/kg	BG8838	ND ^f	ND	ND	37.5	62.5	75.0	62.5	50.0
		WU2	ND	ND	ND	37.5	75.0	87.5	100.0	75.0

^a BG8838 expresses clade 1 PspA, while WU2 and D39 express clade 2 PspA.

^b In these columns, the 4 anti-PspA^{hkR36A} MAbs were the most protective.

^c Surface staining and complement deposition are expressed as fold change (x) in the GMFI value relative to the corresponding isotype control.

^d Bactericidal activity of the anti-PspA^{hkR36A} MAb is represented as percent killing. A higher value indicates higher bactericidal activity. The assay was not performed with strain WU2.

^e Mouse protection is expressed as percent survival. A higher value indicates higher protective efficacy. This experiment was not done with strain D39.

^f ND, not determined.

strains that bear PspA of a different PspA family. We screened our panel to identify the anti-PspA^{hkR36A} MAbs that exhibited the maximum reactivity across PspA clades. Our ELISA and flow cytometry-based surface staining data revealed that all of the 18 anti-PspA MAbs (raised against the clade 2-expressing strain R36A) recognized family 1 PspA and not PspAs representing families 2 and 3. This is consistent with the observation that mice immunized with DNA vaccine expressing the extracellular domain of PspA were protected from strains bearing PspA of the same clade when tested in an intraperitoneal challenge mouse model. Briles and coworkers, however, reported that human anti-

PspA antibodies, when administered to mice, conferred protection against strains expressing PspA belonging to families 1 and 2 (29).

Antibody-dependent complement-mediated phagocytosis is a well-established mechanism of pneumococcal clearance (22). Antibodies directed at pneumococci help in clearance by augmenting opsonization. Antibodies to PspA enhance complement deposition on the pneumococcal surface, thereby contributing to their protective effect (24). A previous report suggested that polyclonal sera against PspA from families 1 and 2 help in enhancing complement deposition (34). Our anti-PspA^{hkR36A} MAbs M4F4, P1E11, P2A4, and P2B5 augmented complement deposition on the 3 PspA family 1-expressing strains analyzed (Table 3). The observed variation in the degree to which various anti-PspA^{hkR36A} MAbs augmented complement deposition across strains may have to do with the chemical nature of the capsule on the target strain, the thickness of the capsule, and the relative accessibility of the pneumococcal surface.

There is evidence to suggest that not all antibodies to PspA are protective indirectly, implying that not all PspA epitopes elicit protective antibody responses. In our previous work, we had observed that P1E11 and P2A4 compete for binding with a PspA^{R36A} subfragment corresponding to a 193- to 286-amino-acid stretch (PspA^{R36A 193-286}), indicating that these two MAbs recognize the same or an overlapping epitope (30). Thus, M4F4, P1E11, P2A4, and P2B5 recognize at least 3 topologically distinct epitopes. The epitopes recognized by P1E11, P2A4, and P2B5 were localized to the PspA^{R36A 193-286} subfragment. The epitope recognized by M4F4 was mapped to the subfragment PspA^{R36A 98-192} (30). Our data are consistent with those reported by Roche et al., who localized the cross protection-eliciting region of PspA to the N-terminal 115 amino acid residues and ~104 C-terminal residues of the extracellular domain from strain EF3296 (12). Knowledge of the epitopes recognized by anti-PspA^{hkR36A} MAbs that do

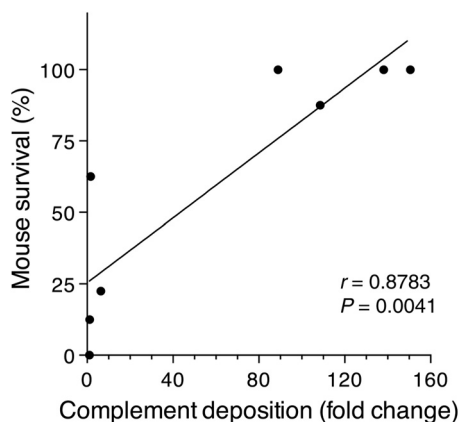


FIG 5 Analysis of 8 anti-PspA^{R36A} MAbs to assess a possible correlation between MAb-dependent complement deposition and their ability to passively protect mice from an otherwise lethal challenge with strain WU2. Each dot represents an anti-PspA^{R36A} MAb. The correlation between *in vitro* complement deposition and passive protection was highly significant by a parametric Pearson's linear regression correlation analysis ($r = 0.8783$, $P = 0.0041$).

and do not elicit protective responses might be put to use to engineer a PspA vaccine that maximizes the proportion of protective antibodies in the antibodies generated. Fine mapping of the conserved B cell epitopes recognized by the protective anti-PspA^{hkR36A} MAbs M4F4, P1E11, P2A4, and P2B5 can help in development of a superior PspA-based vaccine.

Our data indicate that complement deposition on pneumococci can be used as a surrogate for the *in vivo* protection for strains like WU2 (Fig. 5). Roche and coworkers had demonstrated that the antibody titer and surface staining do not correlate with *in vivo* protection and thus are not useful as a surrogate for protection (12). Cohen et al. reported that a whole-cell ELISA is an inadequate predictor of *in vivo* protection (11). While the modified surface killing assay for PspA developed by Genschmer et al. is likely to be significant (9), complement deposition as an *in vitro* correlate of protection is easier to perform and amenable to automation. The complement deposition assay might be potentially useful in quantitating the relative protective efficacy of antibodies against novel protein vaccine antigens (or their subfragments) to confer protection *in vivo*.

ACKNOWLEDGMENTS

This work was supported in part by the intramural research program of the National Institute of Immunology and by grant BT/PR5037/MED/15/77/2012 from the Department of Biotechnology (DBT), India.

Naem Khan was the recipient of a Senior Research Fellowship from DBT.

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