

Monoclonal Antibodies to *Shigella* Lipopolysaccharide Are Useful for Vaccine Production

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There is a significant need for an effective multivalent *Shigella* vaccine that targets the most prevalent serotypes. Most *Shigella* vaccines under development utilize serotype-specific lipopolysaccharides (LPSs) as a major component based on protection and epidemiological data. As vaccine formulations advance from monovalent to multivalent, assays and reagents need to be developed to accurately and reproducibly quantitate the amount of LPSs from multiple serotypes in the final product. To facilitate this effort, we produced 36 hybridomas that secrete monoclonal antibodies (MAbs) against the O antigen on the LPS from *Shigella flexneri* 2a, *Shigella flexneri* 3a, and *Shigella sonnei*. We used six of these monoclonal antibodies for an inhibition enzyme-linked immunosorbent assay (iELISA), measuring LPSs with high sensitivity and specificity. It was also demonstrated that the *Shigella* serotype-specific MAbs were useful for bacterial surface staining detected by flow cytometry. These MAbs are also useful for standardizing the serum bactericidal assay (SBA) for *Shigella*. Functional assays, such as the *in vitro* bactericidal assay, are necessary for vaccine evaluation and may serve as immunological correlates of immunity. An *S. flexneri* 2a-specific monoclonal antibody killed *S. flexneri* 2b isolates, suggesting that *S. flexneri* 2a LPS may induce cross-protection against *S. flexneri* 2b. Overall, the *Shigella* LPS-specific MAbs described have potential utility to the vaccine development community for assessing multivalent vaccine composition and as a reliable control for multiple immunoassays used to assess vaccine potency.

Shigella is a genus of Gram-negative bacteria that is responsible for bacillary dysentery, or shigellosis, which is a major public health problem in the world, particularly in developing countries. It is estimated to cause about 90 million cases of severe diarrhea and about 100,000 deaths per year (1). In developed countries, *Shigella* causes a significant number of cases of diarrhea among children in day care settings, travelers, and military personnel (2). There are four species of *Shigella*: *Shigella dysenteriae*, *Shigella flexneri*, *Shigella boydii*, and *Shigella sonnei*. *S. sonnei* and *S. flexneri* are common throughout the world, but *S. dysenteriae* and *S. boydii* are rare in the United States (3).

A clear and well-defined correlate of immunity has not been fully established to guide the rational design of an effective *Shigella* vaccine (3). Although cross-reactions among different lipopolysaccharides (LPSs) have been reported (4, 5), protection against shigellosis appears to be serotype specific, highlighting the significant role of immune responses to the O-antigen (O-Ag) portion of the LPS (6, 7). Antibodies to O-Ag have been used to define at least 50 serotypes among *Shigella* isolates (3, 8). *S. flexneri* has 14 serologically unique serotypes, which have O-Ags with distinct molecular structures (9). The serotype-specific epitopes (except for type 6) are generated by lysogenic phages that add O-acetyl groups or glucose residues to the common *S. flexneri* O-Ag tetrasaccharide repeating units (3, 9, 10). The O-Ag of *S. sonnei* type I (phase I) has an O-Ag disaccharide repeating unit (8, 11), which is produced by genes in a large (180 to 200 kb) plasmid (pINV) (11, 12). When *S. sonnei* loses pINV, it produces LPSs without O-Ag (phase II) and thus is rough and becomes avirulent (11). The structures of LPS O-Ags from *S. dysenteriae* and *S. boydii* differ from those of *S. flexneri* and *S. sonnei* (9).

Antibodies to *Shigella* O-Ag can fix complement on the target bacteria and kill them in a serotype-specific manner. Thus, in the wake of successful multivalent pneumococcal conjugate vaccines, a leading approach for *Shigella* vaccines is to use LPSs conjugated

to carrier proteins (13–16). LPS serotypes included in the vaccines are chosen based on their geographical prevalence and their induction of cross-reactivity (13–16). LPS conjugate vaccines often include LPSs from *S. flexneri* 2a, *S. flexneri* 3a, and *S. sonnei* since they are commonly isolated in many parts of the world (17). The development of *Shigella* LPS conjugate vaccines can be facilitated if one can easily identify the target bacterial strains for vaccine production, measure the quantity of LPSs in the vaccine lots, and determine vaccine-induced immunity. Thus, we have produced monoclonal antibodies (MAbs) to these three serotypes, and we demonstrate their utility in vaccine development assays, such as inhibition enzyme-linked immunosorbent assay (iELISA), bacterial surface staining by flow cytometry, and bactericidal assay (BCA).

MATERIALS AND METHODS

Bacterial isolates. Eleven reference strains of *Shigella* were obtained from either the ATCC or the Walter Reed Army Institute of Research (WRAIR) in Washington, DC, as shown in Table 1. An additional 49 clinical isolates from Kenya were serotyped at WRAIR and were supplied by WRAIR to the University of Alabama at Birmingham (UAB), which confirmed the

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TABLE 1 Reference strains of *Shigella* used in the study

| Strain and serotype | Reference strain name | Source |
|-------------------------|-----------------------|--------|
| <i>S. dysenteriae</i> 1 | ATCC 9361 | ATCC |
| <i>S. dysenteriae</i> 2 | ATCC 9750 | ATCC |
| <i>S. flexneri</i> 2a | ATCC 700930 | ATCC |
| <i>S. flexneri</i> 2a | 2457T | WRAIR |
| <i>S. flexneri</i> 2b | ATCC 12022 | ATCC |
| <i>S. flexneri</i> 3a | J17B | WRAIR |
| <i>S. flexneri</i> 5a | M90T-Wash | WRAIR |
| <i>S. boydii</i> 1 | ATCC 9207 | ATCC |
| <i>S. sonnei</i> 1 | ATCC 9290 | ATCC |
| <i>S. sonnei</i> 1 | 53G | WRAIR |
| <i>S. sonnei</i> 1 | Moseley | WRAIR |

serotyping results using the *Shigella* antisera set 2 kit (product 294838; Denka Seiken Co. Ltd, Tokyo, Japan).

MAB production. Female BALB/c mice (The Jackson Laboratory, Bar Harbor, ME) at 4 to 6 weeks of age were used for immunization with *S. flexneri* 2a, *S. flexneri* 3a, or *S. sonnei* LPSs. All studies were carried out using a protocol (APN 10019) approved by the Institutional Animal Care and Use Committee of the University of Alabama at Birmingham. The mice were immunized intraperitoneally with 10 µg of LPS/mouse on days 0, 14, and 28. *S. flexneri* 2a and *S. sonnei* LPSs were obtained from the WRAIR and the Lanzhou Institute of Biological Products (LIBP; Lanzhou, China), and *S. flexneri* 3a LPS was from the WRAIR. LPS was extracted using the hot phenol method of Westphal and Jann (18). After the increased level of antibodies to LPSs was confirmed by ELISA on day 35, mice were boosted intraperitoneally with 10 µg of the same LPS/mouse on day 60. The spleen was removed on day 64, and splenocytes were fused with SP2/0-Ag14. Primary culture wells were screened with LPS ELISA (as described below), and ELISA-positive wells were cloned twice by limiting dilution. The MAb isotypes were determined using the IsoStrip mouse monoclonal antibody isotyping kit (product 11493027001; Sigma). Cloned hybridoma cell lines were kept frozen in liquid nitrogen. Hybridoma supernatant was collected after the hybridoma cell was allowed to grow to the terminal stage in the serum-free medium Hybridoma-SFM (product 12045-076; Gibco).

LPS ELISAs. The reactivity of MAbs with LPSs isolated from different *Shigella* serotypes was determined by ELISA. *S. flexneri* 3a LPS was obtained from the WRAIR, and the LPSs of *S. flexneri* 2a and *S. sonnei* were obtained from LIBP. LPS from *Escherichia coli* 0111:B4 was purchased from Sigma Chemical (St. Louis, MO). Each well of an ELISA plate was coated with 100 µl of 1 µg/ml of LPS in phosphate-buffered saline (PBS) for 18 h at 4°C, washed, and blocked with 0.5% bovine serum albumin (BSA) in PBS. Undiluted samples were loaded into each well and incubated for 1 h. After unbound antibodies were washed away, alkaline phosphatase-conjugated anti-mouse Ig (product A0162; Sigma) was added to each well. After a 1-h incubation, the wells were washed and incubated with *p*-nitrophenyl phosphate (pNPP), and the optical densities of the plates were measured at 405 nm (OD₄₀₅). All reactions were performed at room temperature.

For the LPS iELISA, hybridoma supernatants were diluted either 1:20 (Hflex2a1, Hsoni1, and Hsoni5) or 1:40 (Hflex2a4, Hflex3a2, and Hflex3a5) in buffer containing various concentrations of the competing LPS (shown in Fig. 1) and were added to LPS-coated ELISA plates. After incubation for 1 h, unbound antibodies were removed by washing, and the wells were incubated for 1 h with alkaline phosphatase-conjugated anti-mouse Ig (product A0162; Sigma). Then, the wells were washed and incubated with pNPP, and their optical densities were measured at 405 nm.

Bacterial surface staining by flow cytometry. Flow cytometric analysis was performed as previously described (19). In brief, frozen bacterial aliquots were washed and resuspended in a flow cytometry buffer (phosphate-buffered saline [PBS; 140 mM NaCl, 3 mM KCl, 5 mM Na₂HPO₄, 2

mM KH₂PO₄, pH 7.4] with 3% fetal bovine serum and 0.01% sodium azide). Aliquots of 50 µl (containing ~5 × 10⁵ CFU) were incubated with 50 µl of hybridoma culture supernatant (diluted 1:2 to 1:250) at 4°C for 30 min. After incubation, bacteria were washed twice and incubated with goat anti-mouse Ig antibody conjugated with phycoerythrin (PE) (product 550589; BD Biosciences). After being washed, the bacteria were examined in a FACSCalibur flow cytometer (Becton Dickinson, Mountain View, CA). The data were analyzed with FCS Express versions 3.0 and 4.0 (De Novo Software, Los Angeles, CA).

Serotyping by agglutination. The bacterial isolates were serotyped with undiluted hybridoma supernatants as described in the *Shigella* antisera set 2 kit (Denka Seiken Co. Ltd, Tokyo, Japan). Briefly, each strain was grown on a sheep blood agar plate (BAP) overnight at 37°C. A small portion of a bacterial colony was taken from the BAP and mixed with 10 µl of MAb supernatant on a microscope slide. After the slide was gently tilted back and forth for 1 min, agglutination was examined under indirect light. Only clear and strong agglutinations were scored as positive.

Bactericidal assay procedure. The BCA was performed by mixing 20 µl of a diluted hybridoma supernatant, 10 µl of target bacteria (~10⁵ CFU/ml), and 50 µl of 16% baby rabbit serum (product 31061-3; Pel-Freez) as the source of complement in a 96-well microtiter plate. Strains 2457T, J17B, and Moseley were used as target bacteria for *S. flexneri* 2a, *S. flexneri* 3a, and *S. sonnei*, respectively. To simplify the assay, a target bacterial strain was grown in Luria-Bertani (LB) broth and frozen in aliquots at -70°C, and an aliquot was thawed and used for each BCA. After 2 h of incubation at 37°C, 10 µl of reaction mixture was applied to a

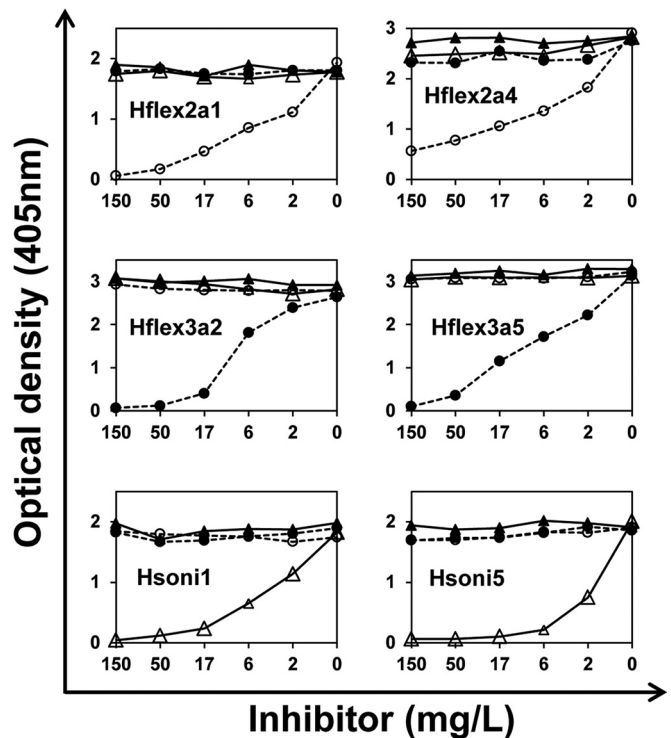


FIG 1 Amount of MAbs bound to the LPS-coated plate (*y* axis) at various inhibitor concentrations (*x* axis) in iELISA. MAbs used for each iELISA are shown in each graph. Inhibitors are LPSs from *S. flexneri* 2a (open circles), *S. flexneri* 3a (solid circles), *S. sonnei* (open triangles), and *E. coli* (solid triangles). Values of 4.7 mg/liter and 5.1 mg/liter of *S. flexneri* 2a LPSs are needed for 50% inhibition of Hflex2a1 and Hflex2a4, respectively. Values of 8.9 mg/liter and 8 mg/liter of *S. flexneri* 3a LPSs are needed to inhibit Hflex3a2 and Hflex3a5, respectively. Values of 3.4 mg/liter and 1.5 mg/liter of *S. sonnei* LPSs are necessary for Hsoni1 and Hsoni5, respectively. All of the experiments were performed more than two times, and one representative example is shown.

TABLE 2 Properties of MAbs to *S. flexneri* 2a, *S. flexneri* 3a, and *S. sonnei*

| Immunizing LPS serotype | MAb name | Isotype | Specificity by ELISA plates coated with LPS from ^a | | | | Specificity by flow cytometry with formalin-fixed target bacteria ^b | | |
|-------------------------|-----------|---------|---|-----------------------|------------------|----------------|--|-----------------------|------------------|
| | | | <i>S. flexneri</i> 2a | <i>S. flexneri</i> 3a | <i>S. sonnei</i> | <i>E. coli</i> | <i>S. flexneri</i> 2a | <i>S. flexneri</i> 3a | <i>S. sonnei</i> |
| <i>S. flexneri</i> 2a | Hflex2a1 | IgG3 | + | – | – | – | + | – | – |
| | Hflex2a2 | IgG2b | + | + | – | – | + | – | – |
| | Hflex2a3 | IgM | + | ± | – | ± | ± | – | – |
| | Hflex2a4 | IgM | + | – | – | – | + | – | – |
| | Hflex2a5 | IgM | + | – | – | – | + | – | – |
| | Hflex2a6 | IgM | + | – | – | – | + | – | – |
| | Hflex2a7 | IgM | + | – | – | – | + | – | – |
| | Hflex2a8 | IgM | + | – | – | – | + | – | – |
| | Hflex2a9 | IgM | + | – | – | – | + | – | – |
| | Hflex2a10 | IgM | + | + | – | – | – | – | – |
| <i>S. flexneri</i> 3a | Hflex3a1 | IgG2b | – | + | – | – | – | + | – |
| | Hflex3a2 | IgG3 | – | + | – | – | – | + | – |
| | Hflex3a3 | IgG2b | + | + | – | – | – | – | – |
| | Hflex3a4 | IgG2b | + | + | – | ± | – | – | – |
| | Hflex3a5 | IgM | – | + | – | – | – | + | – |
| | Hflex3a6 | IgM | – | + | – | – | – | + | – |
| | Hflex3a7 | IgM | – | + | – | – | – | + | – |
| | Hflex3a8 | IgM | – | + | – | – | – | + | – |
| | Hflex3a9 | IgM | – | + | – | – | – | + | – |
| | Hflex3a10 | IgM | – | + | – | – | – | + | – |
| | Hflex3a11 | IgM | – | + | – | – | – | + | – |
| | Hflex3a12 | IgM | – | + | – | – | – | + | – |
| | Hflex3a13 | IgM | – | + | – | – | – | + | – |
| | Hflex3a14 | IgM | – | + | – | – | – | + | – |
| | Hflex3a15 | IgM | – | + | – | – | – | + | – |
| | Hflex3a16 | IgM | – | + | – | – | – | + | – |
| | Hflex3a17 | IgM | – | + | – | – | – | + | – |
| | Hflex3a18 | IgM | – | + | – | – | – | + | – |
| <i>S. sonnei</i> | Hsoni1 | IgG3 | – | – | + | – | + | – | + |
| | Hsoni2 | IgM | – | – | + | – | – | – | + |
| | Hsoni3 | IgM | – | – | + | – | – | – | + |
| | Hsoni4 | IgM | – | – | + | – | – | – | + |
| | Hsoni5 | IgM | – | – | + | – | – | – | + |
| | Hsoni6 | IgM | – | – | + | – | – | – | + |
| | Hsoni7 | IgM | – | – | + | – | – | – | + |
| | Hsoni8 | IgM | – | – | + | – | – | – | + |

^a ELISA results were scored as “–” when the OD₄₀₅ was less than 0.2, “±” when OD₄₀₅ was greater than or equal to 0.2 but less than 0.5, and “+” when OD₄₀₅ was greater than or equal to 0.5. The OD₄₀₅ of wells with no supernatant was ~0.1. Undiluted hybridoma supernatants were used for ELISA.

^b Target strains used for flow cytometric studies were ATCC 700930, J17B, and ATCC 9290, respectively, for *S. flexneri* 2a, *S. flexneri* 3a, and *S. sonnei*. The flow cytometric results are scored as “–” when real signal/background signal (S/N) is less than 2, “±” when S/N is greater than or equal to 2 but less than 5, and “+” when S/N is greater than or equal to 5. Background is the geometric mean fluorescence obtained with no MAb, and the signals are 20, 17, and 16 mean fluorescence units for *S. flexneri* 2a, *S. flexneri* 3a, and *S. sonnei*, respectively. Optimally diluted (ranging from 1:2 to 1:250) supernatants were used for flow cytometry.

small area of a square petri dish (12 cm by 12 cm) containing LB agar. Each petri dish was divided into 48 areas to accommodate reaction mixtures from 48 wells of the microtiter plate. The culture was incubated overnight at 29°C to limit the colony size. The agar plate was then overlaid with agar containing 0.1% NaN₃ and 100 µg/ml of triphenyltetrazolium chloride (TTC), which is reduced by the growing bacteria to a red compound resulting in colored colonies. The petri dish was photographed with a digital camera, and the number of colonies was determined with a computer program called NICE (NIST’s integrated colony enumerator) (20).

RESULTS

Screening MAbs for binding specificity with LPS ELISA and flow cytometry. As shown in Table 2, the cell fusions yielded a large number of hybridoma clones secreting about 1 to 30 µg/ml of

MAbs. To select the hybridoma clones that are useful for further studies, we first determined their immunoglobulin isotypes. Most clones secreting LPS-specific antibodies were IgM, but a few clones were IgG (Table 2). The IgG subclass secreted by the hybridomas was either IgG2 or IgG3, which is consistent with previous reports on MAbs binding T cell-independent type 1 antigens like LPS (21). Antibody-binding specificities of undiluted hybridoma supernatants were also determined by ELISA (summarized in Table 2). As expected, all MAbs bound to the homologous LPS used for immunization. In several cases, the *S. flexneri* MAbs reproducibly showed cross-reactivity with other *S. flexneri* serotypes. For instance, Hflex2a2, Hflex2a3, and Hflex2a10 cross-reacted with *S. flexneri* 3a. Conversely, Hflex3a3

TABLE 3 Slide agglutination test with six selected MAbs^c

| Target bacterial serotype | Strain name | Hflex2a1 or Hflex2a4 | Hflex3a2 or Hflex3a5 | Hsoni1 or Hsoni5 |
|------------------------------------|-------------------------|----------------------|----------------------|------------------|
| <i>S. dysenteriae</i> ^a | MHK03853 | — | — | — |
| | MHK04764 | — | — | — |
| | NTS01572 | — | — | — |
| | MHK04372 | — | — | — |
| <i>S. dysenteriae</i> 1 | ATCC 9361 ^b | — | — | — |
| <i>S. dysenteriae</i> 2 | ATCC 9750 ^b | — | — | — |
| <i>S. flexneri</i> 1a | NTS01435 ^b | — | — | — |
| <i>S. flexneri</i> 1b | MHK03955 ^b | — | — | — |
| <i>S. flexneri</i> 2a | MHK03875 ^b | + | — | — |
| | MHK03876 | + | — | — |
| | MHK04642 | + | — | — |
| | MHK04649 | + | — | — |
| | MHK04687 | + | — | — |
| | MHK04964 | + | — | — |
| | MHK04720 | + | — | — |
| | NTS01277 | + | — | — |
| | NTS01021 | + | — | — |
| | NTS01011 | + | — | — |
| | ATCC 700930 | + | — | — |
| | 2457T ^b | + | — | — |
| <i>S. flexneri</i> 2b | ATCC 12022 ^b | — | — | — |
| <i>S. flexneri</i> 3a | MHK03936 ^b | — | + | — |
| | MHK03563 | — | + | — |
| | MHK05049 | — | + | — |
| | NTS01591 | — | + | — |
| | MHK03620 | — | + | — |
| | MHK04508 | — | + | — |
| | MHK04754 | — | + | — |
| | MHK03792 | — | + | — |
| | MHK01327 | — | + | — |
| | MHK03857 | — | + | — |
| | J17B ^b | — | + | — |
| <i>S. flexneri</i> IV ^a | MHK04643 ^b | — | — | — |
| | MHK04947 | — | — | — |
| | NTS01425 ^b | — | — | — |
| | MHK03739 | — | — | — |
| | MHK04790 | — | — | — |
| <i>S. flexneri</i> 5a | M90T-Wash ^b | — | — | — |
| <i>S. flexneri</i> 6 | MHK04055 | — | — | — |
| | NTS00780 ^b | — | — | — |
| | NTS00731 ^b | — | — | — |
| | NTS01352 | — | — | — |
| | MHK03789 | — | — | — |
| <i>S. boydii</i> ^a | MHK03826 | — | — | — |
| | NTS01627 | — | — | — |
| | MHK03680 | — | — | — |
| <i>S. boydii</i> 1 | ATCC 9207 ^b | — | — | — |
| <i>S. sonnei</i> I | MHK04042 | — | — | + |
| | MHK04536 | — | — | + |
| | NTS01648 ^b | — | — | + |
| | ATCC 9290 | — | — | + |
| | 53G ^b | — | — | + |
| | Moseley | — | — | + |

(Continued on following page)

TABLE 3 (Continued)

| Target bacterial serotype | Strain name | Hflex2a1 or Hflex2a4 | Hflex3a2 or Hflex3a5 | Hsoni1 or Hsoni5 |
|---------------------------|-----------------------|----------------------|----------------------|------------------|
| <i>S. sonnei</i> II | MHK03809 ^b | — | — | — |
| | MHK03868 | — | — | — |
| | NTS00870 ^b | — | — | — |
| | MHK04321 | — | — | — |
| | MHK04955 | — | — | — |
| | NTS01169 | — | — | — |
| | NTS00961 | — | — | — |

^a Subtypes are unknown.

^b These are the 19 strains used for the iELISA studies.

^c +, test results showed large and unambiguous clumps; —, results showed no evidence of clumps.

and Hflex3a4 cross-reacted with *S. flexneri* 2a. Also, Hflex2a3 and Hflex3a4 cross-reacted with *E. coli* LPS, although the cross-reaction was weak.

To independently confirm the specificities of these MAbs, we tested their binding to formalin-fixed *Shigella* bacteria by flow cytometry (Table 2). Most of the flow cytometric data are consistent with the ELISA data, with a few exceptions, which primarily involved cross-reactive antibodies. Hflex3a3 and Hflex3a4 did not bind the target bacteria in flow cytometry. No MAbs against *S. flexneri* LPS cross-reacted with each other or *S. sonnei*, but the *S. sonnei* MAb Hsoni1 cross-reacted with *S. flexneri* 2a. Thus, the observed cross-reactivity may be assay dependent and may not reflect true biologically meaningful cross-reactivity. Taken together, our hybridomas produce MAbs binding the O-Ag and may not bind the more conserved inner core of LPS. Based on these results, we chose 6 out of 36 hybridomas—Hflex2a1, Hflex3a2, Hsoni1, Hflex2a4, Hflex3a5, and Hsoni5—as representative clones for further studies. Two hybridomas (one IgG and one IgM) were chosen for each serotype.

Ability of *Shigella*-specific MAbs to agglutinate clinical isolates and reference strains of *Shigella*. The serotype specificities of the six selected hybridomas were investigated further for their ability to agglutinate clinical isolates representing all four different species of *Shigella* (Table 3). The clinical isolates were first serotyped with the conventional rabbit serotyping reagents. Undiluted supernatants of the six hybridomas expressing either IgG or IgM MAbs induced clumping. Furthermore, MAbs reacted only with the homologous serotypes and produced results identical to those obtained with the rabbit antisera. Thus, hybridomas produce MAbs specific to the O-Ag of LPS.

Studying the six MAbs for binding LPS with iELISA and flow cytometry. A valuable assay for *Shigella* vaccine production would be able to measure the amount of LPSs from different serotypes contained within a multivalent formulation. As an iELISA based on MAbs can be used to measure LPS concentrations, we investigated the six selected hybridomas for an iELISA. As seen in Fig. 1, iELISA can be developed using hybridoma supernatants diluted 1:20 (Hflex2a1, Hsoni1, and Hsoni5) or 1:40 (Hflex2a4, Hflex3a2, and Hflex3a5), and the iELISA using the MAbs was sensitive (down to 1 to 10 mg/liter) and highly specific to the homologous serotype.

To investigate the usefulness of iELISA in serotyping *Shigella* isolates, we examined culture supernatants from 19 selected strains (indicated in Table 3) with iELISA based on MAbs Hflex2a4, Hflex3a5, and Hsoni5. The selected strains represent the

four species of *Shigella*. Supernatants of strains expressing LPS homologous specificity of iELISA inhibited more than 40% at 1:3 dilutions; however, supernatants from strains expressing nonhomologous serotypes inhibited 20% or less at 1:3 dilutions (see Fig. S1 in the supplemental material). The findings suggest that *S. flexneri* 2a, *S. flexneri* 3a, and *S. sonnei* lysates have about 11 μg/ml, 38 μg/ml, and 5 μg/ml of LPS, respectively. Thus, the iELISA is also useful in typing bacterial isolates.

Inactivated whole bacteria are often investigated as *Shigella* vaccine candidates (22). To investigate whether MAbs from the six hybridomas can be used to characterize such whole-cell vaccines, we studied MAbs for their usefulness in the staining of individual bacteria using flow cytometry. MAbs demonstrated selective and unambiguous staining of bacteria when the MAbs were used at optimal dilutions (described in Fig. 2). Generally, IgM antibodies required higher dilutions than IgG antibodies, perhaps because IgM antibodies may clump the bacteria and prevent staining at lower dilutions.

Bactericidal activity of *Shigella*-specific MAbs. An important measure of vaccine-induced immunity is studying immune serum for its ability to kill bacteria *in vitro*. Since MAbs can be useful as controls or standards in bactericidal assays, we examined the supernatants from the six selected hybridomas for killing various *Shigella* strains. To obtain supernatants useful for BCA, hybridoma cells were grown in an antibiotic-free medium without animal sera (Hybridoma-SFM; Gibco) since animal sera often have bactericidal activity (data not shown). Monoclonal antibody Hflex2a1 killed *S. flexneri* 2a isolates specifically (Fig. 3), but Hflex2a4 killed *S. flexneri* 2b isolates in addition to *S. flexneri* 2a isolates (Fig. 3). Consistent with this observation, Hflex2a4 also bound *S. flexneri* 2b by flow cytometry (Fig. 2). Hflex3a2 and Hflex3a5 killed *S. flexneri* 3a isolates only, and Hsoni1 and Hsoni5 killed *S. sonnei* isolates only (Fig. 3). Interestingly, some survivors were noted with target strain Moseley (*S. sonnei*). The survivor may have lost the pINV plasmid spontaneously and escaped killing by antibodies. Also, we noted that high concentrations of IgM antibodies showed a prozone effect in killing (see Fig. S2 in the supplemental material). Thus, BCAs with IgM MAbs were performed after a predilution of 1:100 to 1:5,000. Interestingly, the MAbs that showed a cross-reaction with *S. flexneri* 2a LPS by ELISA (e.g., Hflex3a3 and Hflex3a4) did not cross-kill *S. flexneri* 2a isolates (see Fig. S3 in the supplemental material). None of our MAbs killed *S. flexneri* 5a, *S. dysenteriae* 1, *S. dysenteriae* 2, or *S. boydii* 1 target bacteria (see Fig. S4 in the supplemental material).

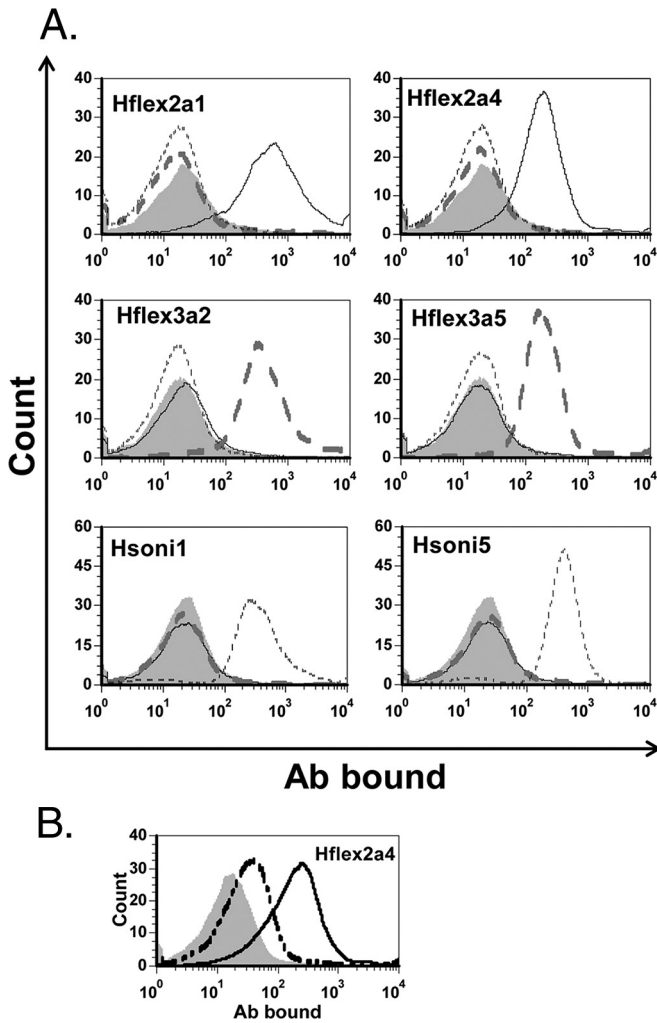


FIG 2 Flow cytometry results with different target bacteria and MABs, which are identified in each graph. (A) The hybridoma supernatants were diluted 1:2 (Hflex2a1), 1:50 (Hsoni1, Hsoni5), or 1:250 (all others). Target bacteria are *S. flexneri* 2a (black solid line), *S. flexneri* 3a (gray dashed line), and *S. sonnei* (black dotted line). Gray filled areas show negative controls obtained with the bacteria (targeted by the MAB) in the absence of MAB. (B) Flow cytometric results with *S. flexneri* 2b target bacteria stained with Hflex2a4 diluted 1:10 (solid line) or 1:250 (dashed line) and with no MAB (gray filled area). All of the experiments were performed more than two times, and one representative example is shown here.

DISCUSSION

Previous studies have described MABs to LPSs from *S. flexneri* 2a (23, 24), *S. flexneri* 3a (25), and *S. sonnei* (11, 26). However, the MABs were not characterized for the assays, such as BCA and flow cytometry, that are useful for vaccine development. Herein, we describe the production of MABs to three common serotypes of *Shigella* and demonstrate their usefulness in iELISA, bacterial surface labeling by flow cytometry, and BCA, which are important assays for *Shigella* vaccine development. As the MABs are serotype specific, they likely target the O-Ag and not the more-conserved inner core of the LPS.

We found iELISA based on these MABs to be very useful for measuring the concentrations of LPS or O-Ag in bacterial culture supernatants or the purified LPS that is used to make conjugate

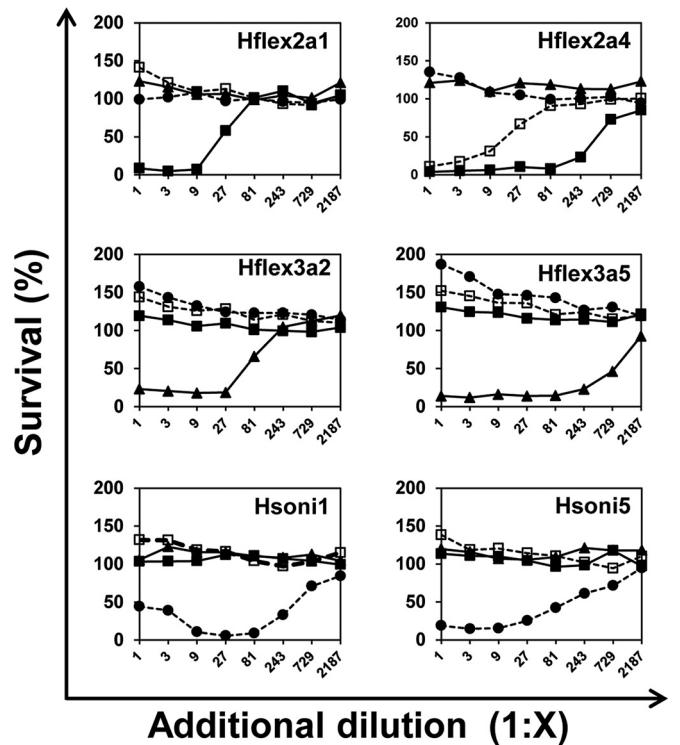


FIG 3 Bactericidal assay results. The relative numbers of surviving target bacteria (y axis) at various additional dilutions of a MAB (x axis) are shown. Names of the MABs are indicated in each graph. Serial dilution was performed for each MAB after a predilution, which was 1:5,000 for Hflex3a5 against target strain *S. flexneri* 3a; 1:1,000 for Hflex2a4 against target strains *S. flexneri* 2a, *S. flexneri* 2b, and *S. flexneri* 3a; or 1:100 for Hflex3a2 against *S. flexneri* 3a and for Hsoni5 against *S. sonnei*. Target strains were 2457T for *S. flexneri* 2a (solid square), ATCC 1202 for *S. flexneri* 2b (open square), J17B for *S. flexneri* 3a (solid triangle), and ATCC 9290 for *S. sonnei* (solid circle). All of the experiments were performed more than two times, and one representative example is shown here.

vaccines. At the moment, LPS concentrations are determined by chemical assays or bioassays that do not distinguish different serotypes of LPSs. For instance, the WRAIR uses Endosafe-PTS, manufactured by Charles River (R. W. Kaminski, personal communication). Inhibition ELISA is a very simple, efficient, and reliable assay that can be established only with hybridoma supernatant without the need to purify MABs. Inhibition ELISA is also sensitive, as it can detect 1 to 10 mg/liter of LPS (Fig. 1). Inhibition ELISA is also highly specific. It can be used to determine the serotype and, with appropriate MABs, also detect O-acetylation, which is immunologically important yet could be lost during O-Ag purification for vaccine production. In contrast, the O-acetylation may be difficult to monitor with bioassay or chemical assays. Moreover, although the conventional chemical assays or bioassays cannot be multiplexed, MAB-based iELISA can be easily multiplexed with bead array technology (e.g., Luminex), as we have shown with pneumococcal antibodies (27). Since future LPS conjugate vaccines will be multivalent, the MAB-based iELISAs should be very useful in vaccine development by measuring LPS concentrations or acetylations of multiple types of LPS during vaccine production.

Immunoassays for *Shigella* LPSs would be useful for nonvaccine studies as well. For instance, the multiplexed iELISA may be

very useful for serotyping *Shigella* isolates as was done for pneumococci (27). Another important application may be a rapid, direct, and simple detection of LPSs in stool samples with dipstick assays for shigella LPS (24, 26). Such an assay would obviate stool culture and facilitate the epidemiologic studies necessary for developing LPS conjugate vaccines in resource-poor locations.

Flow cytometric assays have been infrequently used with bacterial targets until recently. With the maturation of instrumentation, flow cytometry permits one to easily examine the expression of LPSs or other virulence factors at a single-cell level. Flow cytometric assays with these MAbs may be used to monitor the loss of pINV and consequently O-Ag from *S. sonnei* cultures in fermentors. It can be used to study the mixtures of target bacteria, which will be used to make multivalent whole-cell vaccines. Also, we can simultaneously monitor other parameters. For instance, one can simultaneously monitor the expression of LPS as well as some other protein virulence factors that may elicit protective antibodies. We believe that flow cytometric assays should be generally useful in whole-cell vaccine development.

BCAs should play an important role in *Shigella* vaccine evaluations because of the lack of animal models that replicate human infections (development of gastrointestinal symptoms with low doses of infection) (4, 5). An important observation was that these MAbs can kill target bacteria in an *in vitro* BCA, and therefore, these MAbs can be useful in the additional development of BCAs and/or for their standardization. Interestingly, the BCA against *S. sonnei* showed some survivors (very small percentage) when Moseley was used as the target. While these survivors were too few to affect BCA results, the survivors may have lost the plasmid pINV and produce LPSs with no O-Ag.

Induction of cross-protection is an important consideration in designing multivalent *Shigella* vaccines. Although previous studies have demonstrated that vaccination with *S. flexneri* 2a or with a recombinant vaccine expressing *S. flexneri* 2a O-Ag generates cross-reactive antibodies in a variety of species (4, 5), these studies used binding assays or animal models that do not replicate human diseases. Carlin and Lindberg reported that ELISA cross-reactivity was not reproduced with agglutination assays (25). Also, we show that cross-reactivity of MAbs can be demonstrated with one assay and not the other (i.e., ELISA versus flow cytometry). Thus, cross-binding may not imply actual cross-protection. In our studies, Hflex2a4 was shown to bind both *S. flexneri* 2a and *S. flexneri* 2b by flow cytometry and to kill both *S. flexneri* 2a and *S. flexneri* 2b target bacteria. Perhaps, the *S. flexneri* 2a LPS may elicit antibodies that not only cross-react with the *S. flexneri* 2b LPS but also cross-protect against *S. flexneri* 2b.

Taken together, our hybridomas should be useful in *Shigella* vaccine development. Our MAbs can serve as standards for these important assays for vaccine development, and they can be used for epidemiologic studies. To make our MAbs available to the public and enhance their broad use, we plan to deposit the six selected hybridoma lines in a cell-line bank.

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