Protection and Long-Lived Immunity Induced by the ID93/GLA-SE Vaccine Candidate against a Clinical *Mycobacterium tuberculosis* Isolate

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*Mycobacterium tuberculosis* HN878 represents a virulent clinical strain from the W–Beijing family, which has been tested in small animal models in order to study its virulence and its induction of host immune responses following infection. This isolate causes death and extensive lung pathology in infected C57BL/6 mice, whereas lab-adapted strains, such as *M. tuberculosis* H37Rv, do not. The use of this clinically relevant isolate of *M. tuberculosis* increases the possibilities of assessing the long-lived efficacy of tuberculosis vaccines in a relatively inexpensive small animal model. This model will also allow for the use of knock-out mouse strains to critically examine key immunological factors responsible for long-lived, vaccine-induced immunity in addition to vaccine-mediated prevention of pulmonary immunopathology. In this study, we show that the ID93/glucopyranosyl lipid adjuvant (GLA)-stable emulsion (SE) tuberculosis vaccine candidate, currently in human clinical trials, is able to elicit protection against *M. tuberculosis* HN878 by reducing the bacterial burden in the lung and spleen and by preventing the extensive lung pathology induced by this pathogen in C57BL/6 mice.

The development of an effective *Mycobacterium tuberculosis* vaccine, and increased availability of novel drugs targeting drug-resistant strains, would significantly contribute to decreasing the continuing spread of this global disease. Nine million people were estimated to have contracted tuberculosis (TB) in 2013, with 1.5 million deaths occurring annually (1). Overall, 3.5% of new cases and 20.5% of previously treated cases are multidrug-resistant (MDR) TB (1), although this can be as high as 35% and 75%, respectively, in central Asian countries. The Beijing lineage of TB has been associated with a number of MDR TB outbreaks (2–6) and accounts for approximately 13% of isolates worldwide (7). The Beijing lineage itself has expanded more rapidly than other lineages (8); this may be due to the higher virulence of modern strains compared with phylogenetically older sublineages (9–11). The clinical isolate, HN878, is a representative strain from the W–Beijing lineage. The HN878 isolate has been studied in mice to characterize the infectivity and pathogenicity of this virulent clinical strain of *M. tuberculosis* (12–14). These same studies have provided insight into the role of the immune response generated to this pathogen in animals and humans and how the immune response to HN878 contributes to the pathology induced following infection with this isolate. Mice infected with the hypervirulent HN878 strain have increased type I interferon (IFN) expression in the lung and decreased Th1 immunity (12), and HN878 infection in IFN-α/β-deficient mice results in decreased bacterial growth (13). In humans, a type I IFN-α molecular signature, expressed by a population of isolated neutrophils, was observed in patients with active TB (15). More recently, analysis using a whole-genome microarray approach showed a similar type I interferon molecular signature in neutrophil-depleted blood from TB-infected patients (16). Ordway et al. have shown a transient increase in IFN-γ from CD4+ T cells in the lung following HN878 infection in mice, but these IFN-γ-secreting cells in the lung were decreased by 30 days, while CD4+ Foxp3+ regulatory T (Treg) cells continued to increase to greater than 30% in the lungs 60 days later (14). Vaccination with *Mycobacterium bovis* bacille Calmette-Guérin (BCG), while effective at 30 days, was shown in the same study to only delay the expansion of the Treg response; Tregs were observed at 60 days at high percentages in the lung at this time point following infection with HN878 (14). Extensive pathology was also observed in the lungs of untreated HN878-infected mice, which survived less than 90 days following infection, whereas pathology was delayed in BCG-vaccinated mice. While survival was prolonged in BCG-vaccinated mice, mice ultimately succumbed to infection (14). This has led Ordway et al. (14) to hypothesize that the inflammation induced by the virulent *M. tuberculosis* strains, including HN878, leads to the induction of CD4+ Foxp3+ Treg cells, followed by suppression or interference with protective host immunity.

Currently, the only TB vaccine approved for humans is *Mycobacterium bovis* bacille Calmette-Guérin (BCG), a live attenuated vaccine dating to the 1920s. BCG has proved effective for preventing severe disseminated disease in children but does not protect against pulmonary TB in adults (17). Additionally, the live attenuated BCG vaccine is unsafe for administration to HIV-pos-
itive or other immunocompromised individuals due to the possibility of developing regional BCG infection (BCG-itis) (18) or disseminated BCG (BCG-osis) (19-22). Development of an effective TB vaccine will be critical for reducing the worldwide burden of this pathogen. Ideally, a new vaccine would provide immunity against growing as well as dormant phases of the bacterium (23) and would also be safe for use in immunocompromised recipients, providing an alternative to the live attenuated BCG vaccine. We have developed a vaccine candidate fusion of 4 M. tuberculosis proteins (Rv1813, Rv2608, Rv3619, and Rv3620, belonging to the PE/PPE [proteins containing Pro-Glu/Pro-Pro-Glu motifs, respectively], EsX, and latency protein categories) (24), designated ID93. Protein antigens used in vaccines frequently result in low immunogenicity (25, 26); thus, we elected to combine ID93 with a Toll-like receptor 4 (TLR4) agonist as an adjuvant.

The TLR4 agonist in our vaccine adjuvant, glupyranosyl lipid adjuvant (GLA), is a synthetic hexacylglipid A analogue. When formulated in an oil-in-water stable emulsion (SE), GLA induces in vivo innate immune responses as well as Th1-skewed cellular immunity to coadministered vaccine antigens (27, 28). GLA adjuvant formulations have been used with experimental vaccines for influenza, HIV (29), respiratory syncytial virus (RSV) (30), leishmaniasis (31), malaria (32), and leprosy (33) in addition to TB (34, 35). Many of these vaccines using GLA formulations have also resulted in protection in preclinical animal models, including mice (34–36), ferrets (37, 38), guinea pigs (34, 35, 39), cotton rats (30), and hamsters (40), against infectious challenge.

Prior studies have shown that mice immunized with the ID93/ GLA-SE candidate vaccine generate a Th1-biased response, with antigen-specific polyfunctional CD4+ T cells. ID93/GLA-SE prophylactic immunization has led to protection in mice against challenge with H37Rv (34, 35) and a multidrug-resistant TN9001 strain of M. tuberculosis (35). When ID93/GLA-SE is used alone as a vaccine, or as a booster to BCG, postinfection survival of guinea pigs is increased (35). Additionally, this vaccine is safe and immunogenic in nonhuman primates (35). Therapeutic vaccination with ID93/GLA-SE has also proved to be an effective adjunct to antibiotic therapy in infected mice and nonhuman primates (41).

Most preclinical efficacy studies of candidate TB vaccines use the M. tuberculosis laboratory strain H37Rv or Erdman. M. tuberculosis H37Rv was isolated in 1905 and was maintained as in vitro cultures until at least 1923 (42); the Erdman strain was isolated in 1945 (43). These strains have not been observed in humans for 70 years or more and, thus, may not be relevant to the M. tuberculosis strains circulating today. Use of clinically relevant M. tuberculosis isolates for preclinical vaccine efficacy and basic research has been recommended (44). M. tuberculosis HN878 is a recent clinical isolate of the Beijing lineage. In this study, we tested the efficacy of ID93/GLA-SE against the W-Beijing strain, M. tuberculosis HN878, using a homologous vaccination strategy; however, for clinical use, this vaccine candidate, and other new TB vaccine candidates, will likely be used as a booster in BCG-vaccinated populations (45, 46).

Here, we tested our hypothesis that ID93/GLA-SE can elicit protection against M. tuberculosis HN878 through reduction of bacterial burden, decreased lung pathology, and increased survival by induction of long-lived Th1 immunity.
deliver 50 to 100 CFU per mouse. Twenty-four hours postinfection, the entire lung was harvested, homogenized, and plated on 7H10 agar plates. Plates were incubated for 3 weeks at 37°C, 5% CO2. Colonies were counted, and the aerosol dose yielding approximately 100 CFU per mouse was selected.

**Aerosol challenge.** Four weeks after the final immunization, mice were challenged with a low-dose aerosol of *M. tuberculosis* HN878. An aerosol exposure chamber (University of Wisconsin, Madison, WI) was calibrated to deliver 50 to 100 viable *M. tuberculosis* HN878 CFU into the lungs (see supplemental material). An aliquot of frozen stock was thawed and diluted in PBS with 0.05% Tween 80 (Sigma-Aldrich). Twenty-four hours after infection, lungs of 3 mice were homogenized and plated onto Middlebrook 7H10 agar (Molecular Toxicology, Inc., Boone, NC) to ensure bacterial delivery of 50 to 100 CFU. Plates were incubated for 2 to 3 weeks at 37°C, 5% CO2, and then examined for colonies. Infection and all subsequent procedures were performed under biosafety level 3 conditions.

**Bacterial burden.** Four weeks after infection with *M. tuberculosis* HN878, 7 mice per group were euthanized with CO2. The accessory lobe of the lung was reserved for histology. Lung and spleen were harvested and homogenized in PBS with 0.05% Tween 80 (Sigma-Aldrich), using an Omni tissue homogenizer (Omni International, Kennesaw, GA). Serial 5-fold dilutions of homogenates were made in PBS with 0.05% Tween 80, and aliquots of the dilutions were plated on Middlebrook 7H10 agar plates. Plates were incubated for 2 to 3 weeks at 37°C, 5% CO2, before counting colonies. Bacterial burden in CFU per organ was calculated and expressed as log10. The reduction in bacterial burden was calculated as mean log10 CFU saline − mean log10 CFU vaccine.

**Survival.** Ten mice from each group were monitored for survival after infection. Moribund animals, and those losing 20% of body weight, were euthanized with CO2. Bacterial burdens in lungs and spleens were determined as described above. The accessory lobe of the lung was reserved for histology. At experiment termination, all remaining mice were euthanized for sample processing in the same manner.

**Histology.** The accessory lobe of the lung was fixed in 10% formalin for at least 7 days. Fixed tissues were embedded in paraffin, cut, and stained with hematoxylin and eosin (H&E) by the Benaroya Research Institute Histology Core Facility (Seattle, WA). Histological analysis was done in a blind manner by group and performed by a board-certified veterinary anatomic pathologist (Brendan Podell, CSU, Fort Collins). Disease severity, based on area affected, cellular composition, and presence of inflammatory tissue injury, was quantitatively and subjectively evaluated in three tissue sections each, from 7 and 10 mice per group at after 4 weeks of infection and at survival endpoints of the study, respectively. Area morphometry for quantification of lesion-to-tissue area proportions was performed on an Eclipse 80i microscope (Nikon Instruments, Melville, NY) equipped with an automated, computer-controlled stage and Stereo investigator software version 11.01 (MBF Bioscience, Williston, VT), with tissue and lesion area estimated using the fraction-fractionator method. Fifteen to 26 counting frames for lungs of mice sampled at week 4 of infection and 22 to 42 counting frames for lungs of mice sampled at study termination were assigned randomly by the software depending on tissue size, and a counting frame of 1,195 μm2 with a grid spacing of 100 μm was used to define the areas of interest. Data were expressed as a percentage ratio of lesion to total tissue area.

**Statistical analysis.** Flow cytometry data were analyzed with FlowJo version 9.7.5 (Treestar, Ashland, OR) and SPICE (National Institutes of Health, http://exon.niaid.nih.gov/spice) using the Wilcoxon signed-rank test. CFU data were analyzed with GraphPad Prism 6 (GraphPad Software, San Diego, CA) using a standard one-way analysis of variance (ANOVA) followed by Tukey’s multiple-comparison test versus saline. A P value of <0.05 was considered significant. Survival curves were compared to saline using the Mantel-Cox log rank test with Bonferroni correction for multiple comparisons, based on a significance level of P < 0.05.

**RESULTS**

ID93 adjuvanted with GLA-SE induces a CD4+ Th1 antigen-specific population. We first conducted an experiment to characterize the immunogenicity of ID93/GLA-SE. Four weeks after the third immunization, splenocytes were harvested from 4 mice and stimulated with ID93. Lymphocytes were gated on CD44+, and cytokine expression in CD4+ or CD8+ T cells was measured. Background expression in unstimulated (medium only) samples was subtracted. Only the ID93/GLA-SE-immunized animals had significant percent frequency of CD4 T cells expressing CD154 (1.5%), IFN-γ, and TNF (1.5%) (see Fig. S1 in the supplemental material). A large majority (>1% frequency) of the activated ID93-specific CD4+ T cells were multifunctional, expressing double cytokines (CD154+ IFN-γ/TNF). Less than 0.1% induction of CD154, IFN-γ, or TNF-expressing CD8+ T cells was observed (data not shown), and no ID93-specific IL-10, IL-17, or IL-5 was produced from CD4+ T cells (see Fig. S1 in the supplemental material). These results are consistent with our previous reports on ID93/GLA-SE immunization (34, 35).

Antigen-specific Th1 cells infiltrate the lungs of ID93/GLA-SE-immunized mice post-HN878 infection. Four weeks after challenge with *M. tuberculosis* HN878, single cell suspensions were prepared from lungs of 3 mice for each group, stimulated with ID93, and analyzed as for preinfection splenocytes for cytokine induction. CD4+ CD44+ T cells of ID93/GLA-SE-immunized mice produced significant amounts of IFN-γ (~2.3%) and TNF (~4%) in response to ID93; significant expression of CD154 (an activation marker) was also induced (Fig. 1A). Small but statistically significant IL-17 induction was generated as well (0.35%) (Fig. 1B). Whereas IL-17 has been shown to be dispensable against infection with *M. tuberculosis* H37Rv, a requirement for IL-17 in protection against HN878 has recently been described (47). CD154+ IFN-γ/TNF double-cytokine-positive cells comprised the majority of the Th1 cytokine-producing CD44+ CD44+ cells, although some CD154+ TNF-positive single positives were seen (Fig. 1C).

ID93/GLA-SE immunization reduces *M. tuberculosis* HN878 bacterial burden. ID93/GLA-SE vaccine efficacy was tested in mice challenged with *M. tuberculosis* HN878. Bacterial titration was performed to determine the dose of the aerosol to include for infection (see Fig. S2 in the supplemental material). BCG-immunized mice were included as a positive control. Four weeks after challenge, organs from 7 mice per group were harvested, and bacterial burdens in lungs and spleens were determined (Fig. 2; see Table S1 in the supplemental material). Compared to saline-injected mice, ID93/GLA-SE immunization reduced CFU in the lungs by 0.54 log10 (P < 0.001) and in the spleen by 0.52 log10 (P < 0.01) (Fig. 2; see Table S1 in the supplemental material). The BCG control was also effective, with protection of 0.96 log10 (P < 0.0001) in the lungs and 0.79 log10 (P < 0.0001) in the spleen (Fig. 2; see Table S1 in the supplemental material). The GLA-SE adjuvant alone did not afford protection in either lung (0.28 log10, P > 0.05) or spleen (0.12 log10, P > 0.05) (Fig. 2; see Table S1 in the supplemental material).

ID93/GLA-SE immunization prolongs survival of mice challenged with HN878. We next determined whether the protective effect afforded by ID93/GLA-SE was long lived. After a low-dose HN878 aerosol challenge, 10 mice per group were monitored for survival. Saline-treated animals had a median survival time of 231 days compared to saline control (Fig. 2C). Median survival times of mice receiving ID93 and ID93/GLA-SE were 41 and 51 days, respectively. The median survival time of mice receiving saline control was 23 days (Fig. 2C).
days, as did those receiving GLA-SE without antigen. Weights were also monitored following infection and were represented as an average for each group over time (Fig. 3A). By study termination on day 248, 80% of mice immunized with ID93/GLA-SE and 90% of BCG-vaccinated controls had survived (Fig. 3B). Both groups were statistically significantly different from saline controls by the Mantel-Cox log rank test with Bonferroni’s correction for multiple comparisons, whereas GLA-SE alone did not enhance survival (Fig. 3B).

FIG 1 ID93-specific Th1 T cells infiltrate the lungs of ID93/GLA-SE immunized mice postchallenge. C57BL/6 mice were immunized 3 times, at 3-week intervals. Four weeks after the final immunization, mice were challenged with low-dose aerosol M. tuberculosis strain HN878. Four weeks later, single-cell suspensions were made from lungs of 3 mice per group and stimulated with ID93 or medium alone. Medium-only background was subtracted from ID93-stimulated samples. (A) Percentage of single-cytokine-producing CD44⁺ CD41001⁻ T cells; (B) percentage of IL-17-producing CD44⁺ CD41001⁺ T cells; (C) Percentage of CD44⁺ CD41001⁺ polyfunctional Th1-cytokine-producing T cells. Bars are mean plus standard deviation; dots indicate individual values. *, P < 0.05 versus saline immunized, using the Wilcoxon signed-rank test.

FIG 2 Bacterial counts within the lungs (A) and spleens (B) of immunized C57BL/6 mice (7 mice per group) 4 weeks after aerosol infection with M. tuberculosis HN878. Results are presented as log10 CFU within each organ. Asterisks represent statistically significant difference from saline using the one-way ANOVA with Tukey’s multiple-comparison test: ***, P < 0.01; ****, P < 0.001; ***** , P < 0.0001.
Longer survival time trends with lower bacterial burden. Bacterial burden was also monitored throughout the survival study. CFU plotted by day of euthanasia yielded a line with a negative slope for the lung (Fig. 4), whereas a negative slope was observed for the spleen (data not shown). Log_{10} CFU data from individual mice were also organized by treatment and time of euthanasia or death and were included as well (Table I).

Decreased lung immunopathology with ID93/GLA-SE in HN878-infected mice. Because infection with M. tuberculosis HN878 has been shown to induce severe pathology in the lungs of untreated mice, we wanted to determine whether ID93/GLA-SE could provide protection against pulmonary immunopathology induced by this M. tuberculosis strain. H&E-stained slides of lung sections from the week 4 time point (Fig. 5) and the survival arm of the study (Fig. 6) were blind and examined by a board-certified veterinary pathologist. At week 4, saline and GLA-SE (adjuvant only) control mice had similar disease profiles, with lesion burdens ranging from 12% to 36% (Fig. 7). Inflammation was mostly composed of macrophages, with few infiltrating neutrophils and lymphocytes (Fig. 5A and C). In contrast, inflammatory cell populations in the lungs of BCG- and ID93/GLA-SE-immunized mice consisted mostly of lymphocytes, with fewer macrophages (Fig. 5B and D). The ID93/GLA-SE and BCG groups showed significantly less lesion involvement and more evidence of protection within the lungs 4 weeks following infection with HN878, ranging from a 7.6% to a 27% lesion-to-tissue burden (Fig. 7). The cellular composition in the lung of BCG-vaccinated mice was similar to that seen in ID93/GLA-SE-immunized mice.

In the survival arm of the study, the trends were similar to week 4 results, with severe disease burden most evident in the untreated saline control group. Saline and GLA-SE adjuvant-alone groups showed evidence of destructive pulmonary inflammation, with high numbers of neutrophils, foamy macrophages, and areas of necrosis with lipid accumulation (Fig. 6A and C). In contrast, ID93/GLA-SE-immunized mice displayed very little pathology following infection with M. tuberculosis HN878, with generalized absence of destructive inflammation and higher proportions of infiltrating lymphocytes with few to absent neutrophils (Fig. 6D). Similar to the observation 4 weeks after infection with M. tuberculosis HN878, the ID93/GLA-SE group showed significantly less lesion involvement and more evidence of protection within the lungs at the terminal time point (8 months following infection), ranging from a 21.3% to a 53.8% lesion-to-tissue burden (Fig. 8). BCG-vaccinated mice exhibited a cellular composition similar to that of the ID93/GLA-SE group but with an increased area of inflammation, primarily due to large lymphoid follicles, some of which contained high proportions of large lymphocytes with perinuclear clearing of the cytoplasm, morphology that is consistent with plasma cells (Fig. 6B and 8). The lesion involvement in the lungs of BCG-immunized mice ranged between 38.3% and 68.7% and was not significantly different from that of the groups given either saline (44.7% to 84.2%) or GLA-SE (42.7% to 79%) (Fig. 8).

In mice euthanized at the end of the experiment, proliferative changes were also present among epithelial cells that line the alveoli. These changes, consistent with squamous metaplasia of pneumocyte epithelial cells, were present in mice injected with saline or adjuvant alone, affecting 7 of 10 animals in each group (Fig. 9). Notably, epithelial hyperplasia and squamous metaplasia were absent from both BCG- and ID93/GLA-SE-vaccinated mice (see representation of epithelial changes within the M. tuberculosis-infected lung tissue; Fig. 9).

**DISCUSSION**

In this study, we show that ID93/GLA-SE, which generates T helper 1 (Th1) CD4+ T cells, elicits long-lived immunity and prevents pulmonary pathology both early (4 weeks) and late (8 months) following challenge with M. tuberculosis HN878. We observed both a reduction in the bacterial burden in the lungs and increased survival in 80% of ID93/GLA-SE-immunized animals up to 250 days following low-dose aerosol challenge with HN878.
One of the characteristic features of this clinical isolate (HN878) is its hypervirulence; this strain has been shown to grow much faster than other *M. tuberculosis* strains, such as H37Rv and Erdman-KO1 (13), which are typically used in challenge models to measure vaccine efficacy.

One of the properties associated with vaccine efficacy is the prevention of lung pathology associated with *M. tuberculosis* infection. An orchestrated balance of host responses against *M. tuberculosis* is required to induce protection through acquired immunity and granuloma formation in the lung, while preventing pathology that can lead to lung injury and exacerbation of disease (48). Tuberculosis, if untreated, can lead to an accumulation of mycobacterial products in alveoli, followed by increased necrosis and caseation within the lung, bacterial growth, and spread of infection (49). TB lesions involved in healing can result in chronic fibrocaseous disease and extensive pulmonary fibrosis, displacing normal lung tissue (49). We have shown that adjuvant formulation plays a key role in the protective efficacy of a tuberculosis vaccine in the lungs of guinea pigs (34). A main difference which results in C57BL/6 mice following infection with the HN878 clinical isolate compared to the H37Rv strain is the increased lung pathology in mice observed with HN878. We found reduced lung inflammation in ID93/GLA-SE-immunized mice, with lung infiltrates consisting mostly of lymphocytes, in contrast to saline or adjuvant-alone groups, which had extensive neutrophil accumulation at both 4 weeks and 8 months following infection with *M. tuberculosis* HN878. Other researchers have also observed greater numbers of lung lesions, increased lesion size, and greater pathology scores (representing greater lung pathology) in mice 30 days following challenge with *M. tuberculosis* HN878 (13). Vaccine efficacy tested in mice against this clinical isolate can therefore be determined based on additional protective parameters besides reduction in bacterial CFU in the lung and decreased dissemination to other organs. Furthermore, since untreated mice die as a result of low-dose infection with HN878, this provides a model for vaccine durability by looking at survival over time. In our experiments, the untreated C57BL/6 mice challenged with *M. tuberculosis* HN878 lived longer than what has been seen in previously published studies (12, 13), although we did observe similar growth characteristics in the lungs of mice 4 weeks after challenge with the HN878 strain (~6.5 log_{10}) (14). There are some differences between our study and these other studies, however, including the

### TABLE 1

<table>
<thead>
<tr>
<th>Group</th>
<th>Time to death in days (bacterial load in lung, log_{10})</th>
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<tbody>
<tr>
<td>Saline</td>
<td>198 (7.21) 209 (6.90) 231 (6.47) 248 (7.18) 240 (7.25) 160 (7.41) 65 (ND) 240 (7.19) 198 (7.06) 234 (6.59)</td>
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<tr>
<td>BCG</td>
<td>248 (6.50) 248 (6.19) 234 (6.51) 248 (6.06) 248 (6.46) 248 (6.40) 248 (6.70) 248 (6.34) 248 (7.00) 248 (6.97)</td>
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<tr>
<td>GLA-SE</td>
<td>234 (6.59) 234 (6.90) 231 (6.64) 240 (7.60) 234 (6.89) 231 (6.58) 149 (8.27) 209 (7.96) 149 (8.80) 174 (7.20)</td>
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<tr>
<td>ID93/GLA-SE</td>
<td>248 (7.17) 248 (6.06) 174 (7.09) 248 (5.98) 248 (6.46) 248 (6.49) 248 (6.38) 248 (6.47) 248 (6.34) 248 (6.16)</td>
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* a Euthanized due to severe ulcerative dermatitis.
  b ND, not done.
  c Fulfilled criteria for euthanasia on experiment end date.

FIG 5 Lymphocyte infiltration is increased and inflammation is reduced in the lungs 4 weeks after *M. tuberculosis* HN878 infection in mice vaccinated with ID93/GLA-SE. (A, C) Lungs of mice receiving saline or GLA-SE alone are dominated by innate immune cell infiltrates consisting of a high proportion of macrophages and neutrophils (arrows). (B, D) In contrast, BCG- and ID93/GLA-SE-vaccinated mice display similar inflammatory infiltrates with frequent incorporation of lymphocytes (arrows), fewer macrophages, and nearly absent neutrophils. Magnification, ×400.
source of mice used in this study. Our study used C57BL/6 mice purchased from Charles River, whereas Ordway et al. (13, 14) used mice obtained from Jackson Laboratories. Sun et al., who also purchased C57BL/6 mice from Charles River, showed that mice survived longer with the HN878 strain (50). It is unclear if using the same mouse strain purchased from two different vendors could result in such differences in resistance to infection with HN878. Recently, Chang et al. showed significant differences in airway responsiveness by measuring lung resistance in response to different doses of methacholine (a method often used in asthma research) in inbred C57BL/6 mice, depending on where the mice originated (mice purchased from five vendors were tested, including Jackson Laboratories and Charles River Laboratories) (51). The authors suggest that environmental factors (husbandry practices, such as caging systems, food components, type of bedding, etc.), rather than genotypic differences between the B6 substrains, were most likely the cause of observed phenotypic differences, particularly since F1 mice bred and raised in the same environment did not show differences in airway responsiveness among the substrains. Manca et al., who also showed higher death rates in untreated mice with HN878 challenge, used B6D2/F1 mice for their study (12) and also used an aerosol system (Lovelace nebu-
oratory. Culturing of the *M. tuberculosis* strain, including the number of passages, could also result in differences seen in different laboratories and could have resulted in decreased virulence in our laboratory. We included our method for culturing the bacterial stock and for determining the challenge dose used (see supplemental data). Regardless of these differences, the utility of the model is still apparent for testing vaccine efficacy with proper controls, due to the induction of lung pathology, decreased survival observed in untreated infected animals (unlike that seen with laboratory *M. tuberculosis* strains), and reduced costs compared to other animal models, such as the guinea pig, for survival studies. Another benefit of using this model would be to enable determination of the immune requirements of durable vaccine protection using this C57BL/6 challenge model, as several knockout models are generated on the C57BL/6 genetic background. To date, our laboratory has shown through adoptive transfer experiments and use of multiple cytokine and cytokine receptor gene knockout mice that the ID93/GLA-SE candidate vaccine does not require IFN-γ, TNF, or inducible nitric oxide synthase (iNOS) for protection against *M. tuberculosis* H37Rv (52). It is unclear, however, whether IFN-γ, TNF, and iNOS elicited by ID93/GLA-SE immunization are critical to long-lived, durable vaccine protection against this clinical isolate and the pathology induced by HN878; these vaccine-induced efficacy readouts are not feasible using the laboratory-adapted H37Rv strain in C57BL/6 mice, since this laboratory *M. tuberculosis* strain is not lethal and significant pathology in the lung due to infection is not observed. Other investigators have suggested that preclinical challenge with relevant clinical isolates in mice and other animal models might be better predictors of vaccine efficacy (14, 53, 54). The data in this study showing protection against a W-Beijing clinical isolate (HN878) support our other protective efficacy studies with ID93/GLA-SE, which have been performed in many animal models and against both sensitive and multidrug-resistant strains of *M. tuberculosis*. Given that the pathogenesis of *M. tuberculosis* HN878 (and other pathogenic strains) may partially stem from the presence of lipids, such as the phenolic glycolipids that induce type I IFN and block Th1 immunity (55) or may be due to transient Th1 immunity, which is diminished due to an expansion of CD4+ Foxp3 Treg cells in response to infection with virulent strains of *M. tuberculosis* (13), we hypothesize that the use of GLA as an adjuvant to induce innate cytokines, such as IL-12p40 (28), to drive a vaccine-mediated Th1 response is an effective strategy against virulent isolates such as HN878. Another possible strategy would be to test the potential synergistic effects of a CCR4 antagonist(s), known to block Treg function (56) combined with the ID93/GLA-SE vaccine. One of these CCR4 antagonists, mogamulizumab, is currently in clinical trials for various cancers, including adult T-cell leukemia-lymphoma, non-small-cell lung cancer, and advanced solid tumors (www.clinicaltrials.gov) (57). Addition of a CCR4 antagonist combined with the MVA85A tuberculosis vaccine candidate showed enhanced purified protein derivative (PPD)-specific IFN-γ-expressing splenocytes in mice approximately 1 week following immunization (58), which supports the feasibility of this strategy.

Another interesting observation in our study was that a majority of mice (7 of 10 mice) in the survival study from both the saline and adjutant control groups displayed epithelial hyperplasia in the lung tissue, suggesting an association between *M. tuberculosis* infection and the development of neoplasia. The morphology observed histologically among the epithelial cells that line the alveoli is representative of squamous metaplasia of pulmonary pneumocytes, which may represent preneoplastic changes within the lung. Squamous metaplasia was observed only in the lungs of mice chronically infected with *M. tuberculosis* HN878; no hyperplasia was present 4 weeks after infection in any of the groups. Chronic infection is a known risk factor for pulmonary cancer, and a recent review of the literature by Falagas et al. has identified several reports showing associations between TB and cancer in humans (59). Another paper recently showed the development of squamous cell carcinoma in mice chronically infected with *tuberculosis* using *M. tuberculosis* Erdman (60). The authors report evidence for the induction of DNA-damaging reactive oxygen and nitrogen within TB-infected macrophages and upregulation of epiregulin mRNA from interstitial macrophages within the TB lesions in the mice. The authors also suggest that epiregulin, an epidermal growth factor, produced from TB-infected macrophages may be serving as a paracrine growth factor during the preneoplasia stages in their model. In their study, infection with *M. tuberculosis* Erdman in C57BL/6 mice led to the appearance of squamous cell carcinoma in 50% of mice (2 of 4 mice) 7 months following infection and in 80% of mice (8 of 10 mice) 12 weeks after infection. The presence of cancer in the lungs was accelerated in B6.C3H-1st1 mice, which include the sst1--tuberculosis susceptible allele (61); 100% (4 of 4 mice) had evidence of squamous cell metaplasia in lungs 5 months after infection. In our study, we used HN878, a W-Beijing clinical isolate, for infection in C57BL/6 mice. Similar to others, we show that the HN878 isolate induces a greater amount of inflammation in the untreated lungs of C57BL/6 mice compared to what is normally observed following infection with the laboratory *M. tuberculosis* strain H37Rv. In addition, C57BL/6 mice also succumb to infection more rapidly with the HN878 strain compared to the rate observed with H37Rv; while C57BL/6 mice will eventually succumb to infection with H37Rv, they are typically considered relatively resistant to infection with
laboratory strains of *M. tuberculosis* (62). This is the first report, to our knowledge, that shows epithelial hyperplasia and squamous metaplasia in the lung with *M. tuberculosis* HN878 in untreated C57BL/6 mice and protection against epithelial hyperplasia and lung pathology, in addition to a reduction in HN878 bacterial load in the lung and spleen and increased survival with ID93/GLA-SE.

In summary, a model that utilizes *M. tuberculosis* HN878 infection in C57BL/6 mice will be useful for testing the efficacy of prophylactic TB vaccines (with and without a BCG prime immunization) where bacterial reduction and immunopathology in the lung can be assessed, and in the context of an adjunct therapeutic vaccine combined with drug treatment against clinical *M. tuberculosis* isolates. Our future plans will include testing ID93/GLA-SE in these scenarios, will investigate whether ID93/GLA-SE inhibits the elicitation of Foxp3+ T regulatory cells in HN878-infected animals and in addition will characterize other important immunological mechanisms of vaccine-induced protection.

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REFERENCES


