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**Listeria-vectored multi-antigenic tuberculosis vaccine protects C57BL/6 and BALB/c mice and guinea pigs against *Mycobacterium tuberculosis* challenge**

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## Abstract

*Mycobacterium tuberculosis* (Mtb) infects one-third of the world's population and is a leading cause of death from a single infectious agent. New TB vaccines are urgently needed to augment immunity conferred by the current modestly protective BCG vaccine. We have developed live attenuated recombinant *Listeria monocytogenes* (rLm)-vectored TB vaccines expressing five [Mpt64/23.5-EsxH/TB10.4-EsxA/ESAT6-EsxB/CFP10-Ag85B/r30] (rLmMtb5Ag) or nine (additionally EsxN-PPE68-EspA-TB8.4) immunoprotective Mtb antigens (rLmMtb9Ag) and evaluated them for safety, immunogenicity and efficacy as standalone vaccines in two mouse models and an outbred guinea pig model. In immunogenicity studies, rLmMtb5Ag administered subcutaneously induces significantly enhanced antigen-specific CD4+ and CD8+ T-cell responses in C57BL/6 and BALB/c mice, and rLmMtb9Ag induces antigen-specific CD4+ and CD8+ T-cell proliferation in guinea pigs. In efficacy studies, both rLmMtb5Ag and rLmMtb9Ag are safe and protect C57BL/6 and BALB/c mice and guinea pigs against aerosol challenge with highly virulent Mtb. Hence, multi-antigenic rLm vaccines hold promise as new vaccines against TB.

## Introduction

*Mycobacterium tuberculosis* (Mtb), the leading cause of death from a single infectious agent, at least until the COVID-19 pandemic, infects approximately one-third of the world's population and causes active tuberculosis (TB) in approximately 10.4 million people and death in ~1.7 million people annually <sup>1</sup>. The only currently approved TB vaccine, bacille Calmette-Guérin or BCG, has been administered to ~5 billion people on earth; it provides only moderate protection against childhood TB and highly variable protection (0 - 87%) against adult pulmonary TB. Hence new TB vaccines are a very high global public health priority and could play an important role in achieving the World Health Organization (WHO) End TB Strategy target of a 95% reduction in TB deaths and a 90% reduction in TB incidence by 2035 compared to 2015 <sup>2</sup>.

While most people in the world are vaccinated with BCG in infancy, especially in low- and middle-income countries where the incidence of TB is highest, most cases of TB occur in adults. In 2020, 89% of new cases of TB occurred in people over 14 years of age<sup>2</sup>. Thus, an effective TB vaccine is needed to boost the immunity of adolescents and adults. In support of this strategy, an impact and cost-effectiveness study of potential TB vaccines in low- and middle-income countries indicated that a vaccine targeting adolescents and adults could have a greater impact than one targeting infants <sup>3</sup>.

Two types of new TB vaccines are under development – prime vaccines to replace BCG and booster vaccines to enhance the immunity of people previously vaccinated with BCG or in the future vaccinated with replacement vaccines for BCG. Prime vaccines

under development, all comprising mycobacteria, include rBCG30, the first vaccine demonstrated more potent than BCG and the first potential replacement vaccine for BCG to enter human trials <sup>4-7</sup>; rBCG(*mbtB*)30, the first vaccine demonstrated more potent and safer than BCG <sup>8</sup>; MTBVAC, an attenuated Mtb <sup>9</sup>, and VPM-1002, a modified BCG <sup>10</sup>. TB booster vaccines under development, all subunit vaccines, include protein/adjuvant vaccines such as r30 (Antigen 85B), the first defined booster vaccine against TB <sup>4</sup>, H1 (Antigen 85B – ESAT6) <sup>11,12</sup>, H4 (Antigen 85B-TB10.4) <sup>13,14</sup>, M72 <sup>15,16</sup>, and ID93 <sup>17,18</sup>; virus vectored vaccines such as MVA85A <sup>19,20</sup>; and bacterium vectored vaccines including Salmonella-vectored vaccines <sup>21</sup> and Listeria-vectored vaccines, including rLm30 <sup>22</sup> and rLmMtb5Ag <sup>23</sup>, as well as rLmMtb9Ag, described herein.

Targeting an adolescent and adult population for a new TB booster vaccine has practical implications regarding the preclinical development and testing of such a vaccine. Since most people in the world in need of an improved TB vaccine have been vaccinated with BCG in infancy, a new TB booster vaccine should demonstrate efficacy in enhancing the protective immunity conferred by BCG in preclinical animal models of pulmonary TB. But if the intended targets of the new TB booster vaccine are adolescents and adults, in whom immunity from BCG vaccination in infancy has largely waned, then a new TB booster vaccine ideally would additionally demonstrate protective efficacy as a standalone vaccine in pre-clinical animal models of pulmonary TB.

Our candidate TB booster vaccine comprises a live attenuated replicating *Listeria monocytogenes* (Lm) bacterium expressing immunoprotective Mtb antigens. The Lm

vector was chosen in large part because of its ability to induce robust antigen-specific CD4<sup>+</sup> and CD8<sup>+</sup> T cell responses to expressed recombinant antigens – both types of T cell immunity are central to immunoprotection against Mtb. The wild-type parent of this vector, a fast growing, Gram-positive, facultative intracellular bacterium that occasionally infects humans and can cause food-borne disease outbreaks, shares important features of its intracellular lifestyle with Mtb. Like Mtb, upon entry into host cells, which include mononuclear phagocytes, Lm initially resides in a phagosome, a site favoring antigen presentation via class II MHC molecules and the induction of antigen-specific CD4<sup>+</sup> T cells. Subsequently, also in common with Mtb, Lm escapes the phagosome and multiplies in the host cytosol, a site favoring antigen presentation via class I MHC molecules and the induction of antigen-specific CD8<sup>+</sup> T cells. As a result of its capacity to induce long-lived cell-mediated immune responses, genetically attenuated Lm has been developed as a vaccine vector for cancer and infectious diseases <sup>24</sup>.

The specific Lm vector that we employ, Lm  $\Delta actA \Delta inlB prfA^*$  (Lm  $\Delta actA \Delta inlB \Delta uvrAB prfA^*$ ) <sup>22,25</sup> has been attenuated from wild-type Lm and rendered more effective as a vaccine vector via several genetic manipulations <sup>25</sup>. First, *actA*, a gene encoding the cell surface transmembrane protein ActA, which promotes intracellular motility via actin polymerization, has been deleted. ActA deletional mutants are able to grow within the cytosol of infected cells, but are unable to induce cell-to-cell spread, resulting in ~1000-fold attenuation in virulence in mouse models <sup>24</sup>. Second, a deletion of *inlB*, encoding internalin B, a virulence factor that promotes invasion of various mammalian cells

including epithelial cells, endothelial cells and hepatocytes, inhibits Lm uptake into non-phagocytic cells, such as hepatocytes, but not into phagocytic cells, including antigen-presenting cells; hence, in the double deletional  $\Delta actA \Delta inlB$  mutant, off-target toxicity is minimized but not antigen presentation of secreted recombinant antigens in antigen presenting cells<sup>26</sup>. Third, a point mutation (G155S) in the master virulence regulator PrfA that renders it constitutively active, promotes Lm escape into the host cell cytosol, and as a result of upregulated expression of PrfA and PrfA-dependent genes, shows enhanced expression of downstream recombinant proteins<sup>25,27,28</sup>.

Using Lm  $\Delta actA \Delta inlB prfA^*$  as a vaccine vector, we have developed several recombinant Lm-vectored Mtb vaccines (rLm) including rLm30<sup>22</sup>, rLmMtb5Ag (rLm5Ag)<sup>23</sup>, and in this study, rLmMtb9Ag (rLm9Ag). The rLm30 vaccine expresses a single Mtb antigen – the 30-kDa major secretory protein or Antigen 85B (r30 or Ag85B, gene Rv1886); rLm5Ag expresses a fusion protein of 5 Mtb antigens – Mpt64/23.5 (Rv1980c), EsxH/TB10.4 (Rv0288), EsxA/ESAT6 (Rv3875), EsxB/CFP10 (Rv3874) and r30; and rLm9Ag expresses, in addition to the 5 antigens expressed by rLm5Ag, a fusion protein comprising 4 additional Mtb antigens - EspA (Rv3616c), EsxN (Rv1793), PPE68 (Rv3873) and TB8.4 (Rv1174c). These 4 additional proteins were selected on the basis of their capacity, when administered as part of a rLm booster vaccine, to enhance protective immunity against Mtb aerosol challenge in BCG-immunized mice<sup>23</sup>. In previous studies of rLm30 and rLm5Ag, we have shown that i) immunization of mice with BCG has no significant effect on local replication or systemic dissemination, growth, and clearance of rLm30 administered intradermally 12 or 15 weeks later<sup>22,23</sup>; ii)

boosting BCG-primed mice with rLm30 and rLm5Ag enhances Mtb antigen-specific CD4+ and CD8+ T cell-mediated immune responses <sup>22,23</sup> ; and iii) boosting BCG-primed C57BL/6 and BALB/c mice with rLm30 and rLm5Ag enhances protective immunity against aerosolized Mtb <sup>22,23</sup>.

Herein, we investigate the immunogenicity and efficacy of rLm5Ag and rLm9Ag as standalone vaccines in three animal models of pulmonary TB – inbred C57BL/6 and BALB/c mice and outbred Hartley guinea pigs. We test them as standalone vaccines - not because we envision them as replacement vaccines for BCG but because, as noted above, most of the people in the world in need of a TB booster vaccine were vaccinated with BCG in infancy; hence, their BCG-induced immunity will have largely waned by the time they would receive a TB booster vaccine many years and often decades later. As testing the potency of a heterologous booster vaccine administered decades after a prime vaccine is not feasible in small animal models, we elected instead to test the vaccines as standalone vaccines so as to mimic the situation in which BCG-induced immunity has completely waned. In so doing, we examined two different strains of mice, C57BL/6 and BALB/c, because these mice display different innate and acquired immune responses to infection, including mycobacterial infection with BCG <sup>29,30</sup>. We additionally examined guinea pigs because these animals develop disease more akin to that of humans than do most strains of mice; e.g., they are highly susceptible to clinical disease after low dose infection with Mtb; they show strong cutaneous delayed-type hypersensitivity to tuberculin; and they display Langhans giant cells in lung lesions and develop caseating granulomas <sup>31</sup>.



In C57BL/6 and BALB/c mice, we show that homologous priming-boosting with rLm5Ag and rLm9Ag vaccines induces antigen-specific CD4+ and CD8+ T cell immune responses and protective immunity against aerosol challenge with virulent Mtb. In guinea pigs, we show that homologous priming-boosting with rLm9Ag induces Mtb antigen-specific lymphocyte proliferation and elevated frequencies of CD8+ T cells in the lungs and/or spleens, and that immunization with rLm5Ag or rLm9Ag induces significant protective immunity against Mtb aerosol challenge.

## Results

### Construction and verification of rLm9Ag expressing fusion proteins of Mtb5Ag and Mtb5AgII from the *comK* and *tRNA<sup>arg</sup>* loci, respectively

To construct rLm5Ag (expressing Mtb Mpt64-EsxH-EsxA-EsxB-r30, with an N-terminal fusion to ActAN, the N-terminal 100 amino acids of Lm ActA)<sup>23</sup> (Supplementary Table 1) and rLm5AgII (expressing ActAN-Mpt64-EsxN-PPE68-EspA-TB8.4) (Supplementary Table 1), we integrated the pPL2e-ActAN-Mtb5Ag and pPL2e-ActAN-Mtb5AgII into the *tRNA<sup>arg</sup>* locus of the rLm chromosome (Fig. 1a, top and middle vaccines). To construct the rLm9Ag vaccine candidate (Supplementary Table 1), we integrated pPL1-ActAN-Mtb5Ag at the *comK* locus followed by integrating pPL2e-ActAN5AgII into the *tRNA<sup>arg</sup>* locus of the Lm chromosome (Fig. 1a, bottom vaccine). This strategy had two advantages. First, the use of two expression cassettes allowed expression of smaller fusion proteins than a single cassette expressing all 9 antigens (estimated Mw of 9-antigen fusion protein of 212 kDa vs. the estimated molecular weights of the 5Ag and 5AgII fusion proteins of 94 and 118 kDa, respectively). The smaller Mw favors better stability and expression. Second, the strategy allows use of the same leader protein of Mpt64 for both the 5AgI and 5AgII expression cassette. We have tested various combinations of Mtb fusion proteins and found that Mtb proteins fused to the C-terminus of Mpt64 tend to be expressed more abundantly than otherwise; hence Mpt64 is evidently an ideal leader protein for the expression cassette of Mtb fusion proteins. We have shown that rLm5Ag, rLm5AgII, and two clones (clones 1 and 3) of rLm9Ag express the expected fusion proteins when grown in broth medium and in infected murine macrophage-like cells. As shown in Fig. 1b and Supplementary Figure 1 left

panels, rLm9Ag vaccine clones 1 and 3 (top panel, lanes 5, 6) express both the 94-kDa 5Ag and the 118-kDa 5AgII fusion proteins (indicated by arrows), similar to the proteins expressed by rLm5AgII (lane 4) and rLm5Ag (lanes 2, 3), respectively, that are detected by a polyclonal antibody to an N-terminal peptide comprising 18 amino acids (A30-K47) of ActA (AK18, courtesy of Justin Skoble and Peter Lauer). As expected, the 94-kDa, but not the 118-kDa, protein band was also detected by the polyclonal antibody to EsxH (middle panel, lanes 2, 3, 5, and 6) which is present in the 5Ag but not the 5AgII fusion protein.

To analyze expression of the heterologous Mtb fusion proteins by rLm growing inside of macrophages, we infected murine macrophage-like cells (J774.A1) with LmVector or rLm expressing various Mtb fusion proteins at a Multiplicity of Infection (MOI) of 10. At 5.5 hours post infection, we harvested the infected cells and analyzed the lysates for protein expression by SDS-PAGE and Western blotting using the polyclonal antibody AK18, which detected the predicted 94-kDa (5Ag) protein band (and non-specific protein bands) from J774A.1 cells infected with rLm5Ag, rLm5AgII, or rLm9Ag clones #1 and 3 (Fig. 1c upper panel and Supplementary Figure 1 right upper panel, lanes 2, 3, 5, and 6, respectively) and the predicted 112-kDa (5AgII) protein band from J774A.1 cells infected with rLm5Ag I and rLm9Ag clones #1 and 3 (Fig. 1c, upper panel, lanes 4, 5, and 6, respectively), but not from LmVector-infected (upper panel, lane 1) and mock infected (upper panel, lane 7) J774A.1 cells. Antibody to Lm P60 detected a ~ 60-kDa protein band from Lm-infected (Fig. 1c lower panel and Supplementary Figure 1 right lower panel, lanes 1 – 6), but not from uninfected cells (Fig. 1c, lower panel, lane 7).

Antibody to  $\beta$ -actin detected a ~42 kDa band from uninfected and infected J774 cell lysates, as expected (Fig. 1c lower panel, and Supplementary Figure 1 right lower panel, lanes 1 – 7).

### Genetic stability and growth kinetics of rLm5Ag and rLm9Ag vaccines

To evaluate the antigen expression cassette stability of rLm grown in broth culture and in infected macrophage-like cells, we examined the growth of rLm vaccine candidates in Brain Heart Infusion (BHI) broth supplemented with various antibiotics and in infected monolayers of the J774A.1 cells with stationary grown rLm vaccines and assayed bacterial replication. As shown in Supplementary Figure 2, the rLm9Ag, rLm5Ag, and rLm5Ag I vaccine candidates grew similarly in the presence or absence of antibiotic selection, either after direct inoculation into BHI broth medium (Supplementary Figure 2a) or after passage in murine macrophage-like cells followed by plating onto BHI agar plates supplemented with various antibiotics (Supplementary Figure 2b & 2c). We verified the heterologous and homologous protein expression by Western blotting as shown in Fig. 1b and Fig. 1c. In addition, we passaged LmVector, rLm5Ag and rLm9Ag vaccines daily for 10 consecutive days on BHI agar plates and verified the stability of the antigen expression cassettes. We also verified the stability of the Mtb antigen expression cassettes after the rLm9Ag vaccine was passaged in guinea pigs (Supplementary Figure 3). Thus, the antigen expression cassettes for Mtb 5Ag (Mpt64-EsxH-EsxA-EsxB-r30), Mtb 5AgII (Mpt64-EsxN-PPE68-EspA-TB8.4), and Mtb 9Ag (5Ag + 5AgII) are stably maintained in the rLm vaccine candidates.

To examine the growth kinetics of rLm vaccine candidates in murine macrophages, we infected monolayers of J774A.1 cells with LmVector or rLm vaccines at an MOI of 10, as described in the legend to Supplementary Figure 2d. In general, all rLm5Ag, rLm5AgI and rLm9Ag vaccines grew similarly to LmVector at 2 h post infection; at 4 h post infection, there were some delays in the growth of rLm5AgII and rLm9Ag clone #1 compared with LmVector; at 6 hours post infection, there were delays in the growth of rLm5AgII and rLm9Ag clone #1 and clone #3; rLm5Ag grew similarly to LmVector at 2, 4, and 6 h post infection (Supplementary Figure 2d). These results suggest that the fusion protein expression cassettes for Mtb 5AgII imposed a modest burden on its bacterial host that caused some growth delay in macrophages.

### rLm5Ag induces antigen-specific T-cell mediated immune responses in mice

We examined the capacity of rLm5Ag to induce antigen-specific T cells and cytokine-expressing CD4<sup>+</sup> and CD8<sup>+</sup> T cells in the lungs and spleens of C57BL/6 mice and BALB/c mice (Fig. 2-5, Supplementary Figure 4 - 7). C57BL/6 mice immunized with rLm5Ag or LmVector produced comparable frequencies of CD3<sup>+</sup> T cells among gated lymphocytes and comparable frequencies of CD4<sup>+</sup>, CD8<sup>+</sup> and CD4-CD8<sup>-</sup> T cells among CD3<sup>+</sup> T cells after 22 h antigen stimulation in the lungs, and for the most part in the spleens, although differences in response to a few antigens were statistically significant (Supplementary Figure 5).

With respect to antigen-specific cytokine-expressing T cells, the rLm5Ag-immunized C57BL/6 mice produced significantly greater frequencies of cytokine-producing CD4<sup>+</sup> T

cells in their lungs and spleens expressing IFN- $\gamma$ , TNF- $\alpha$ , and/or IL-2 and polyfunctional CD4<sup>+</sup> T cells expressing two or more cytokines among IFN- $\gamma$ , TNF- $\alpha$ , and IL-2 in response to *in vitro* 6 h stimulation with Ag85B (Fig. 2e, 2m, 3a, 3e), 5Ag pool (Fig. 2f, 2n, 3b, 3f), or PPD (Fig. 2g, 2o, 3c, 3g) than mice immunized with LmVector. No significant differences in the frequencies of CD4<sup>+</sup> T cells expressing any of the cytokines were detected after *in vitro* stimulation with 23.5/Mpt64 (Fig. 2a, 2i), TB10.4/EsxH (Fig. 2b, 2j), ESAT6/EsxA (Fig. 2c, 2k), or CFP10/EsxB (Fig. 2d, 2l), or as expected PMA (Fig. 2h, 2p). Notably, mice immunized with rLm5Ag also produced greater frequencies of CD8<sup>+</sup> T cells expressing IFN- $\gamma$  and TNF- $\alpha$  and polyfunctional CD8<sup>+</sup> T cells expressing both cytokines in response to *in vitro* stimulation with 23.5/Mpt64 (Fig. 4a, 5a), TB10.4/EsxH (Fig. 4b, 5b), 5Ag pool (Fig. 4f, 5c), and PPD (Fig. 4g, 5d) than mice immunized with LmVector in the lung, and a greater frequency of CD8<sup>+</sup> T cells expressing TNF- $\alpha$  in response to the 5Ag pool in the spleen (Fig. 4n). Of note, the frequencies of antigen specific cytokine-producing CD4<sup>+</sup> and CD8<sup>+</sup> T cells in the lungs and spleens after 22-hour antigen stimulation were similar to those after 6-hour stimulation. Thus, homologous priming-boosting C57BL/6 mice with rLm5Ag induces Mtb antigen-specific cytokine-expressing CD4<sup>+</sup> and CD8<sup>+</sup> T cells, especially Ag85B-specific CD4<sup>+</sup> T cells expressing IFN- $\gamma$ , TNF- $\alpha$  and IL-2, and Mpt64 and EsxH-specific CD8<sup>+</sup> T cells expressing IFN- $\gamma$  and TNF- $\alpha$ .

BALB/c mice immunized three times with rLm5Ag produced substantially greater (~2-5-fold) frequencies of live lung CD3<sup>+</sup> T cells than LmVector-immunized mice after 22h incubation whether incubated with or without antigen ( $P < 0.05$  -  $P < 0.001$ )

(Supplementary Figure 6a); of these greatly expanded numbers of CD3<sup>+</sup> T cells, CD4<sup>+</sup> T cell frequencies were somewhat reduced and CD8<sup>+</sup> and CD4<sup>-</sup>CD8<sup>-</sup> T cell frequencies were somewhat increased, sometimes significantly so, whether incubated with or without antigen ( $P < 0.001$  –  $P < 0.0001$  without antigen) (Supplementary Figure 6b, c, d). In contrast, in the spleen, frequencies of CD3<sup>+</sup>, CD4<sup>+</sup>, CD8<sup>+</sup>, and CD4<sup>-</sup>CD8<sup>-</sup> T cells were generally comparable (Supplementary Figure 6e-h).

With respect to antigen-specific cytokine-expressing CD4<sup>+</sup> T cells, in the spleens, BALB/c mice immunized with rLm5Ag produced significantly greater amounts of IFN- $\gamma$ , TNF- $\alpha$ , and/or IL-17A expressing CD4<sup>+</sup> T cells in response to 6h *in vitro* stimulation with Mpt64, EsxH, EsxB, Ag85B, PPD and GI-H37RV than mice immunized with LmVector; the only antigens not inducing a significantly greater response was EsxA and, as expected, PMA (Supplementary Figure 7, top two rows, CD4<sup>+</sup> T cells). In contrast, with respect to antigen-specific cytokine-expressing CD8<sup>+</sup> T cells, BALB/c mice immunized with rLm5Ag produced moderately greater frequencies of splenic CD8<sup>+</sup> T cells expressing only IL-17A or IL-17A and IFN- $\gamma$  in response to only EsxH and EsxA but not to Mpt64, EsxB, Ag85B, PPD, GI-H37RV and, as expected, PMA (Supplementary Figure 7, bottom two rows, CD8<sup>+</sup> T cells).

Thus, homologous priming-boosting BALB/c mice with rLm5Ag induces Mtb antigen-specific cytokine-expressing CD4<sup>+</sup> and CD8<sup>+</sup> T cells, where the CD4<sup>+</sup> T cells show specificity to Mpt64, EsxH, EsxB, and Ag85B – all but EsxA – and express

predominantly IFN- $\gamma$  and TNF- $\alpha$ , and CD8<sup>+</sup> T cells show specificity to EsxH and EsxA, but express predominantly IL-17A and IFN- $\gamma$ .

Comparing C57BL/6 and BALB/c mice, immunization with rLm5Ag substantially increases the frequencies of live CD3<sup>+</sup> T cells in the lungs of BALB/c mice (Supplementary Figure 6a) but not C57BL/6 mice (Supplementary Figure 5a) independent of antigen. Both mice produce antigen-specific cytokine-expressing CD4<sup>+</sup> and CD8<sup>+</sup> T cells, although the antigen specificity and suite of cytokines secreted differ between these two mouse strains.

### rLm5Ag and rLm9Ag induce protective immunity against aerosol challenge with virulent Mtb Erdman strain in BALB/c and C57BL/6 mice

In preliminary studies, we evaluated the protective efficacy in C57BL/6 and BALB/c mice of a combination of two rLm vaccines expressing 5 antigens - r30/Ag85B, 23.5/Mpt64, TB10.4/EsxH, ESAT6/EsxA, and CFP10/EsxB (combination of rLm30 + rLm4Ag, designated as rLm5Ag\*). First, we explored i.d. and intranasal (i.n.) administration of the composite vaccine rLm5Ag\* in C57BL/6 mice. In these experiments, we used BCG as a positive control against which to compare the efficacy of the Lm vaccines, as BCG consistently provides strong efficacy in animal models of TB. We immunized groups of C57BL/6 mice, 8 per group, with PBS or BCG i.d. or i.n. at Week 0, or the rLm5Ag\* i.d. or i.n. three times at Weeks 0, 7, and 10, challenged all the mice with aerosolized Mtb (average of 24 CFU of the Mtb Erdman strain delivered to the lungs of each mouse, as assayed on Day 1 post-challenge) at Week 13 and



euthanized the mice at Week 23 (Fig. 6a, top panel). Mice immunized i.d. three times with rLm5Ag\* had significantly lower CFUs in their lungs and spleens than Sham-immunized mice ( $P<0.05$  and  $P<0.01$ , respectively), not significantly different from mice immunized i.d. once with BCG (Fig. 6a, left middle and bottom panels); mice immunized i.n. once with BCG or three times with rLm5Ag\* showed no reduction in bacillus burden in the lungs compared with Sham-immunized mice (Fig. 6a, right middle panel), but had significantly lower CFUs in their spleens than Sham-immunized mice ( $P<0.0001$  and  $P<0.05$ , respectively) (Fig. 6a, right bottom panel).

Next, we immunized groups of 8 BALB/c mice with PBS (Sham) or BCG (positive control) i.d. at Week 0, or i.m. with the rLm5Ag\* three times at Weeks 10, 14, and 18 or twice at Weeks 14 and 18, then challenged the mice at Week 22 with aerosolized Mtb (average of 30 CFU of the Mtb Erdman strain delivered to the lungs of each mouse, as assayed on Day 1 post-challenge), and euthanized the mice 10 weeks later (Week 32) to assay bacillus burden in their lungs and spleens (Fig. 6b, upper panel). As shown in Fig. 6b, middle and bottom panels, mice immunized i.m. twice or three times with rLm5Ag\* had significantly lower CFUs in their lungs and spleens than Sham-immunized mice; the reduction in CFU in the lungs was comparable to that of mice immunized with BCG. Three immunizations were slightly more efficacious than two immunizations and resulted in a greater statistically significant difference from sham in both the lung and spleen.

Subsequently, we performed definitive studies comparing the protective efficacy of rLm30, rLm5Ag\* (combination of rLm30 + rLm4Ag), rLm5Ag (single vaccine expressing the same 5 antigens as rLm5Ag\* from the *tRNA<sup>arg</sup>* locus), and rLm9Ag (clones #1 and #3) as a standalone vaccine in both C57BL/6 and BALB/c mice immunized three times s.q., a route that was found to be both practical and efficacious against Mtb aerosol challenge in heterologous prime-boost studies involving a BCG prime and an rLm boost. We immunized groups of 8 C57BL/6 and BALB/c mice s.q. three times at Weeks 0, 3, and 6 with rLm30, rLm5Ag\*, rLm5Ag, or two individual clones of rLm9Ag (rLm9Ag #1 and rLm9Ag #3); unimmunized (UI) mice or mice immunized i.d. with BCG, or three times s.q. with LmVector served as controls. Four weeks after the last immunization, mice were challenged with aerosolized Mtb (average of 10 CFU of Mtb Erdman strain delivered to the lungs of each mouse, as assayed at Day 1 post challenge). At Week 20, the mice were euthanized and organ bacillus burdens assayed (Fig. 7a and 8a).

As shown in Fig. 7, C57BL/6 mice immunized with BCG once or immunized three times with rLm30, rLm5Ag\*, rLm5Ag, and both clone #1 and #3 of rLm9Ag, had lower CFUs in their lungs and spleens than sham-immunized mice and mice immunized with LmVector. Among the 4 rLm vaccines tested, rLm9Ag provided the greatest protection to C57BL/6 mice against Mtb aerosol challenge. Specifically, in the lung, mice immunized with both clones #1 and #3 of rLm9Ag had significantly lower CFU than sham-immunized mice (-0.4 and -0.5 logs, respectively ( $P < 0.05$ )) and mice immunized with LmVector ( $P < 0.01$ ), comparable to mice immunized with BCG (Fig. 7b, left panel). In the spleen, mice immunized with clone #1 of rLm9Ag had significantly lower CFU (-

0.7 log) then sham-immunized mice ( $P<0.05$ ). Mice immunized with clone #3 of rLm9Ag had significantly lower CFU than both sham-immunized mice (-0.8 logs) ( $P<0.05$ ) and mice immunized with LmVector ( $P<0.05$ ), similar to and not statistically significantly different from mice immunized with BCG (Fig. 7b, right panel). These results indicate that vaccinating C57BL/6 mice with rLm vaccines comprising 5 antigens or 9 antigens as a standalone vaccine provides potent protection against aerosol challenge with virulent Mtb Erdman strain and the rLm9Ag vaccine provides protection comparable to BCG.

As shown in Fig. 8, BALB/c mice immunized with BCG once or immunized three times with rLm30, rLm5Ag\*, rLm5Ag, or both clones #1 and #3 of rLm9Ag had lower CFUs in their lungs and spleens than unimmunized mice and mice immunized with LmVector. Among the 5 rLm vaccine candidates tested, rLm5Ag\* (combination of rLm30+rLm4Ag) and rLm5Ag (single vaccine) provided the best protection to BALB/c mice against Mtb aerosol challenge. Specifically, in the lung, mice immunized with rLm5Ag\* and rLm5Ag had significantly lower CFU than sham-immunized mice (-1.1 and -1.1 logs, respectively) ( $P<0.0001$  and  $P<0.0001$ ) and mice immunized with LmVector ( $P<0.0001$  and  $P<0.01$ ), comparable (difference not statistically significant) to mice immunized with BCG. Mice immunized with rLm9Ag #1 and rLm9Ag #3 also had significantly lower CFU than sham-immunized mice (-0.8 and -0.9 logs, respectively) and mice immunized with LmVector ( $P<0.01$  and  $P<0.001$ ) in the lung (Fig. 8b, left panel). In the spleen, mice immunized with rLm5Ag\* and rLm5Ag had significantly lower CFU than sham-immunized mice (-1.6 and -1.7 logs, respectively) ( $P<0.0001$  and  $P<0.0001$ ) and

LmVector (-1.1 and -1.2 logs, respectively) ( $P < 0.001$  and  $P < 0.0001$ ). Mice immunized with rLm9Ag clone #1 and #3 also had significantly lower CFU than unimmunized mice (-0.9 and -1.1 logs, respectively) ( $P < 0.0001$  and  $P < 0.0001$ ) and mice immunized with LmVector (-0.3 and -0.6 logs, respectively) ( $P < 0.001$  and  $P < 0.0001$ ) (Fig. 8b, right panel). These results indicate that in BALB/c mice, vaccinating with rLm5Ag\*, rLm5Ag, or rLm9Ag as a standalone vaccine provides potent protection against aerosol challenge with virulent Mtb Erdman strain.

### rLm9Ag induces antigen specific T-cell proliferation in guinea pigs

To evaluate the capacity of the rLm9Ag vaccine to induce T-cell mediated immune responses in guinea pigs, we immunized guinea pigs with rLm9Ag or LmVector three times at Weeks 0, 3, and 6 and 6 days later evaluated responses of spleen and lung lymphocytes, CD4+ T cells, and CD8+ T cells to stimulation with Mtb peptide antigens. As shown in Fig. 9, in general, lung (Fig. 9a) and spleen (Fig. 9b) cells of guinea pigs immunized with rLm9Ag had greater frequencies of lymphocytes than guinea pigs immunized with LmVector after antigen stimulation. Specifically, in the lungs, guinea pigs immunized with rLm9Ag produced greater frequencies of lymphocytes in response to Ag85B, EspA, EsxA, EsxB, EsxH, EsxN, PE68, TB8.4 and PPD (Fig. 9a). In the spleens, guinea pigs immunized with rLm9Ag produced greater frequencies of lymphocytes in response to Ag85B, EsxA, EsxB, and TB8.4 (Fig. 9b). As expected, there were no significant differences between animals immunized with LmVector and rLm9Ag in frequencies of lymphocytes stimulated without antigen (Medium alone) or with ConA (positive control) (Fig. 9a, 9b). With respect to proliferating CD4+ T cells

(CD4<sup>+</sup> CTV<sup>Low</sup>) in the lung, as shown in Fig. 9c, no significant differences were detected between guinea pigs immunized with LmVector and rLm9Ag after *in vitro* stimulation with Mtb peptide antigens. With respect to proliferating CD4<sup>+</sup> T cells (CD4<sup>+</sup> CTV<sup>Low</sup>) in the spleen, as shown in Fig. 9d, guinea pigs immunized with rLm9Ag had greater frequencies of proliferating CD4<sup>+</sup> T cells to Ag85B and EsxB than guinea pigs immunized with LmVector. With respect to proliferating CD8<sup>+</sup> T cells (CD8<sup>+</sup> CTV<sup>Low</sup>) in the lung, as shown in Fig. 9e, there were no significant differences between guinea pigs immunized with LmVector and rLm9Ag after *in vitro* stimulation with Mtb peptide antigens. With respect to proliferating CD8<sup>+</sup> T cells (CD8<sup>+</sup> CTV<sup>Low</sup>) in the spleen, as shown in Fig. 9f, guinea pigs immunized with rLm9Ag had greater frequencies of CD8<sup>+</sup> T cells in response to Ag85B, EsxB, EsxN, PPE68, TB8.4, and PPD than guinea pigs immunized with LmVector.

Thus, rLm9Ag induces significantly increased lung and/or spleen cells in response to 8 of its 9 recombinant Mtb antigens, significantly increased proliferating splenic CD4<sup>+</sup> T cells in response to Ag85B and EsxB, and significantly increased proliferating CD8<sup>+</sup> T cells in response to 5 of its 9 Mtb antigens.

### rLm5Ag and rLm9Ag induce protective immunity against aerosol challenge with virulent Mtb Erdman strain in outbred guinea pigs

Finally, we evaluated the capacity of the rLm5Ag and rLm9Ag vaccines to induce protective immunity against Mtb aerosol challenge in the outbred guinea pig model. As shown in Fig. 10a, we immunized guinea pigs three times s.q. at Weeks 0, 3, and 6 with

the rLm multi-antigenic vaccines or LmVector, challenged them at Week 10 with aerosolized Mtb Erdman strain, subsequently monitored their weight for 10 weeks, and then euthanized them to determine CFU in their lungs and spleens; guinea pigs immunized with BCG i.d. served as a positive control. As shown in Fig. 10b, Supplementary Figure 8 and Supplementary Table 2, guinea pigs immunized three times at Weeks 0, 3, and 6 with rLm9Ag at low, medium, and high doses, or with rLm5Ag at a medium dose gained significantly more weight 4-10 weeks post-challenge than guinea pigs immunized with LmVector ( $P < 0.05$  –  $P < 0.0001$ ); the weight gain of rLm vaccinated guinea pigs was not significantly different from that of guinea pigs immunized with BCG (except guinea pigs immunized with the low dose of rLm9Ag at Week 10 post challenge). Similarly, as shown in Fig. 10c, guinea pigs immunized with low, medium, or high doses of rLm9Ag had lower Mtb bacillus burdens than guinea pigs immunized with LmVector in their lungs (0.2, 0.8, and 0.4 logs lower, respectively) (Fig. 10c, left panel) and spleens (0.2, 1.2 and 0.9 logs lower, respectively) (Fig. 10c right panel). Similarly, guinea pigs immunized with rLm5Ag also had lower bacterial burdens than LmVector immunized guinea pigs in the lung and spleen (0.8 and 0.9 logs lower, respectively), reductions virtually equivalent to those in animals immunized with the medium dose of rLm9Ag. Guinea pigs immunized with BCG also had significantly lower Mtb bacillus burdens in their lungs (Fig. 10c left panel) (1.4 logs lower) and spleens (Fig. 10c right panel) (2.2 logs lower) than those immunized with LmVector. Among the 3 doses of rLm9Ag tested, the medium dose ( $10^6$ ) of rLm9Ag provided the best protection against aerosol challenge with highly virulent Mtb. Thus, both the rLm5Ag and rLm9Ag vaccines induce potent protective immunity in guinea pigs.

## Discussion

Our study shows that homologous priming-boosting of inbred C57BL/6 and BALB/c mice and outbred guinea pigs with rLm5Ag (expressing a fusion protein of Mtb antigens Mpt64-EsxH-EsxA-EsxB-Ag85B) and rLm9Ag (additionally expressing Mtb antigens Mpt64-EsxN-PPE68-EspA-TB8.4) induces antigen-specific CD4<sup>+</sup> and CD8<sup>+</sup> T-cell mediated immunity and immunoprotection against aerosol challenge with virulent Mtb Erdman in all three animal models.

Both CD4<sup>+</sup> and CD8<sup>+</sup> T cells are required to control primary TB infection. CD4<sup>+</sup> T cells help CD8<sup>+</sup> T cells maintain effector function and prevent exhaustion, and the synergy between CD4<sup>+</sup> and CD8<sup>+</sup> T cells promotes the survival of mice infected with Mtb <sup>32</sup>.

Our results show that rLm5Ag induces both CD4<sup>+</sup> and CD8<sup>+</sup> T cell-mediated immune responses in both C57BL/6 and BALB/c mice. In both mouse models, immunizing with rLm5Ag or rLm9Ag induces elevated frequencies of lung and/or splenic CD8<sup>+</sup> T cells, consistent with Lm's reputation as a potent inducer of CD8<sup>+</sup> T cells <sup>24</sup>. Similarly, in guinea pigs, rLm9Ag induces proliferating antigen-specific CD4<sup>+</sup> and CD8<sup>+</sup> T cells. CD8<sup>+</sup> T cells appear to play a more important role in protection against Mtb in primates than in rodents <sup>33</sup>. Hence, studies in rodents may underestimate the efficacy of an rLm vaccine in non-human primates and humans.

In C57BL/6 mice, rLm5Ag induced significantly elevated frequencies of antigen-specific polyfunctional CD4<sup>+</sup> and CD8<sup>+</sup> T cells expressing IFN- $\gamma$ , TNF- $\alpha$ , and sometimes IL-2 (the CD4<sup>+</sup> T cells in response to stimulation with Ag85B, and the CD8<sup>+</sup> T cells in

response to stimulation with Mpt64 and EsxH); the frequency of IL-17A expressing cells was not significantly elevated. In BALB/c mice, rLm5Ag induced significantly elevated frequencies of CD4<sup>+</sup> T cells expressing IFN- $\gamma$  and TNF- $\alpha$  (in response to stimulation with Mpt64, EsxH, EsxB, and Ag85B), and additionally significantly elevated frequencies of CD4<sup>+</sup> and CD8<sup>+</sup> T cells secreting IL17A (in response to EsxH and EsxA). Hence, these two mouse strains displayed a somewhat different CD4<sup>+</sup> and CD8<sup>+</sup> T cell cytokine expression profile after rLm5Ag immunization.

Among the five antigens common to both rLm5Ag and rLm9Ag, all five induced T cell responses in at least one animal model, and four induced T cell responses in multiple animal models. Antigen 85B was an especially dominant antigen, inducing T cell responses in all three animal models – cytokine-expressing CD4<sup>+</sup> T cells in both C57BL/6 and BALB/c mice and proliferating CD4<sup>+</sup> and CD8<sup>+</sup> T cells in guinea pigs. EsxB also stood out, inducing cytokine-expressing CD4<sup>+</sup> T cells in BALB/c mice and proliferating CD4<sup>+</sup> and CD8<sup>+</sup> T cells in guinea pigs. EsxH induced cytokine-expressing CD4<sup>+</sup> T cells in both C57BL/6 and BALB/c mice. Finally, Mpt64 induced cytokine-expressing CD4<sup>+</sup> T cells in BALB/c mice and CD8<sup>+</sup> T cells in C57BL/6 mice. The rLm9 Ag vaccine, containing 4 additional Mtb antigens, was tested for immunogenicity only in guinea pigs. Three of these four new antigens – EsxN, PPE68, and TB8.4 – induced proliferating CD8<sup>+</sup> T cells in guinea pigs.



An Lm-vectored vaccine has major advantages as a TB vaccine including i) Lm multiplies rapidly intracellularly and secretes foreign antigens into the host cell cytosol, as noted above, and then is rapidly cleared – 7 to 10 days post immunization <sup>22</sup>; ii) Lm-vectored vaccines with  $\Delta actA \Delta inlB$  deletions have an established safety profile in humans; the vaccines were well tolerated in a Phase I study <sup>34</sup>; iii) pre-existing immunity to Lm <sup>35,36</sup> and to BCG does not deleteriously affect immunization with Lm-vectored vaccines <sup>22</sup>, in contrast to some virus- and mycobacterium-vectored vaccines; iv) Lm-vectored vaccines have enhanced capacity to induce both CD4+ and CD8+ T cell-mediated immune responses, and is an especially potent inducer of CD8+ T cells, as noted above; and v) Lm-vectored vaccines can be cheaply manufactured in broth medium at large scale without the need for extensive purification as with protein/adjuvant vaccines or virus-vectored vaccines grown in mammalian cells.

Although in this study we tested the Listeria vectored vaccines as standalone vaccines, we envision an rLm vaccine not as replacement vaccine for BCG but as a heterologous booster vaccine for people previously vaccinated with BCG, or in the future with an improved mycobacterial vaccine that eventually replaces BCG. Greater than 5 billion people on earth who have been vaccinated with BCG live in TB endemic areas, and hence might benefit from a heterologous TB booster vaccine. As most of these people, including adolescents and adults, would have been vaccinated with BCG in infancy, their BCG-induced immunity is likely to have largely waned by the time they were to receive such a booster vaccine many years and decades later. For this reason, as noted earlier, we considered it important to test the efficacy of our Listeria vectored

vaccines as standalone vaccines. As standalone vaccines, we did not expect the rLm vaccines to be superior to BCG, which shares thousands of antigens with Mtb. However, that our studies demonstrate that the rLm vaccines are in many cases comparable in potency to BCG is noteworthy and, in our view, strongly supports the continued development of these rLm vaccines as TB booster vaccines.

In conclusion, in this study, the protective efficacy of multi-antigenic rLm TB vaccines – including rLm5Ag and rLm9Ag – was demonstrated in three rigorous animal models of pulmonary TB. This follows upon the demonstration in previous studies of the protective efficacy of a single-antigen rLm vaccine (rLm30) and multi-antigenic rLm vaccines –including rLm5Ag – as booster vaccines to enhance the level of immune protection afforded by BCG immunization. Hence, an rLm vaccine expressing multiple Mtb immunoprotective antigens has substantial promise as a new vaccine to combat the TB pandemic.

## Material and methods

### Ethics statement

All animals were maintained in a specific-pathogen-free animal facility and used according to protocols approved by the UCLA Institutional Animal Care and Use committee.

### Cell lines, bacteria, animals, protein antigens, and antibodies.

Murine (J774A.1, ATCC TIB-67) monocytes were cultured as we described previously<sup>22</sup>. *M. bovis* BCG Tice and Mtb Erdman (ATCC 35801) strains were acquired and stocks prepared as we described previously<sup>22</sup>. The Listeria vector, Lm  $\Delta actA \Delta inlB prfA^*$ <sup>22,25</sup>, derived from *Listeria monocytogenes* 10403S strain (phage-cured, DP-L4056)<sup>37</sup>, and recombinant Lm-vectored vaccines were grown to mid-log phase in Yeast Extract broth medium, collected by centrifugation, resuspended in PBS, titrated, and stored in 20% glycerol at -80°C until use. Six to eight-week-old female C57BL/6 and BALB/c mice were purchased from Envigo (Indianapolis, IN) or Jackson Laboratory (Bar Harbor, Maine, USA) and three-week-old outbred male Hartley strain guinea pigs were purchased from Charles River Laboratories (Wilmington, MA, USA). The following Mtb protein reagents were obtained through BEI Resources, NIAID, NIH: Ag85B (Gene Rv1886c), Purified Native Protein from Strain H37Rv, NR-14857; ESAT-6, Recombinant Protein Reference Standard, NR-49424; CFP-10, Recombinant Protein Reference Standard, NR-49425; Mpt64, Recombinant Protein Reference Standard, NR-44102; and

GI-H37RV, Mtb, Strain H37Rv, Gamma-Irradiated Whole Cells, NR14819. The Mtb protein EsxH/TB10.4 (gene Rv0288) was obtained from Aeras (formerly Rockville, Maryland, United States). Rabbit polyclonal antibody to ActAN (AK18, lot D4698) was obtained courtesy of Justin Skoble and Pete Lauer; rabbit polyclonal antibody to TB10.4 was obtained from Aeras (formerly Rockville, Maryland, United States); monoclonal antibody to Lm P60 (P6007, Lot AG-20A-0022-C100) was purchased from AdipoGen (San Diego, United States); and monoclonal antibody to  $\beta$ -actin (A5441) was purchased from Sigma (St. Louis, United States).

### Construction of Lm-vectored multi-antigenic vaccines.

We constructed Lm-vectored multi-antigenic rLm vaccine candidates using the *Lm*  $\Delta actA \Delta inlB prfA^*$  vector<sup>25</sup> and two Lm site-specific phage integration vectors, pPL1 and pPL2, through conjugation process, as previously described by us and others<sup>22,23,37</sup>. The pPL1 conjugation vector (kindly provided by Peter Lauer) utilizes the listeriophage U153 integrase and attachment site for insertion at the *comK* locus of the rLm chromosome and carries a Gram-positive chloramphenicol acetyltransferase gene; the pPL2e-derived conjugation vector [pBHE666 containing *actA* promoter and the N-terminal 100 amino acids of ActA (ActAN), kindly provided by Justin Skoble] modified from pPL2, utilizes the listeriophage PSA integrase and attachment site for insertion in the 3' end of the *tRNA<sup>arg</sup>* gene of the rLm chromosome and carries an erythromycin resistance gene<sup>37</sup>. We cloned the genes encoding Mtb proteins, optimized for expression of Mtb proteins in *Listeria monocytogenes* and purchased from DNA2.0 (currently <https://www.atum.bio/>) (Newark, CA), into pPL1 and pBHE666 by the

restriction enzyme method and in some cases by the Electra Vector System (<https://www.atum.bio/>) to generate the following plasmids: pPL2e-ActAN-Mtb4Ag (for integration of ActAN-Mtb4Ag expression cassette at the *tRNA<sup>arg</sup>* locus to construct rLm4Ag), pPL2e-ActAN-Mtb5Ag (for integration of ActAN-Mtb5Ag expression cassette at the *tRNA<sup>arg</sup>* locus to construct rLm5Ag), pPL1-ActAN-Mtb5Ag (for integration of ActAN-Mtb5Ag expression cassette at the *comK* locus to construct rLm9Ag), and pPL2e-ActAN-Mtb5AgII (for integration of Mtb5AgII expression cassette at the *tRNA<sup>arg</sup>* locus to construct rLm9Ag). We have deposited the sequences of these plasmids to Genbank (<https://www.ncbi.nlm.nih.gov/genbank/>). All molecular plasmid constructs were confirmed by restriction enzyme digestion and nucleotide sequencing. Candidate vaccines rLm30 (expressing ActAN-r30/Ag85B) (Supplementary Table 1) <sup>22</sup>, rLm4Ag (expressing the fusion protein ActAN-Mpt64-EsxH-EsxA-EsxB) (Supplementary Table 1), and rLm5Ag (expressing the fusion protein ActAN-Mpt64-EsxH-EsxA-EsxB-r30) (Supplementary Table 1) were constructed previously <sup>23</sup> [where rLm5Ag is referred to as rLm5Ag(30)]; the Mtb protein expression cassettes in these vaccines were cloned into the pPL2e-derived vector and integrated at the *tRNA<sup>arg</sup>* locus of the rLm chromosome. The rLm5AgII vaccine candidate (Supplementary Table 1), expressing the Mtb fusion protein of ActAN-Mpt64-EsxN-PPE68-EspA-TB8.4 from the pPL2e vector integrated at the *tRNA<sup>arg</sup>* locus as well, was constructed similarly as described previously <sup>22,23</sup>. The rLm9Ag vaccine candidate (Supplementary Table 1) was constructed by integrating the pPL1-ActAN-Mtb5Ag (expressing ActAN-Mpt64-EsxH-EsxA-EsxB-r30) at the *comK* locus followed by integrating the pPL2e-ActAN-5AgII (expressing ActAN-Mpt64-EsxN-PPE68-EspA-TB8.4) at the *tRNA<sup>arg</sup>* locus. The resultant rLm9Ag carries a total of 9 Mtb

antigens with Mpt64 being a common antigen located at the N-terminus of both 5Ag and 5AgII fusion proteins. The pPL1-ActAN-Mtb5Ag conjugation vector carries a codon-optimized antigen expression cassette for the fusion protein of Mpt64( $\Delta$ 1V-23A)-RP-EsxH-GGSG-EsxA-GSSGGSSG-EsxB-GSSGGSSG-Ag85B( $\Delta$ 2Q-43A) (abbreviated as Mpt64-EsxH-EsxA-EsxB-r30), in which RP is a dipeptide encoded by an EagI restriction enzyme site, and GSSG and GSSGGSSG are flexible fusion protein linkers. The pPL2e-ActAN-Mtb5AgII conjugation vector carries a codon-optimized antigen expression cassette for the fusion protein of Mpt64( $\Delta$ 1V-23A)-EsxN-GSSG-PPE68-GSSGGSSG-EspA( $\Delta$ 111F-193L)-GSSGGSSG-TB8.4( $\Delta$ 2R-28A) (abbreviated as Mpt64-EsxN-PPE68-EspA-TB8.4).

### Growth kinetics and stability of Lm-vectored multi-antigenic vaccines *in vitro*

The growth kinetics of rLm in broth and in murine macrophage-like cells were evaluated as described previously by us <sup>22,23</sup>. To assay the stability of rLm vaccines grown on agar plates, we passaged the Lm vector, rLm5Ag and rLm9Ag daily for 10 consecutive days on BHI agar plates supplemented with streptomycin, and at day 5 and day 10, we transferred 20-25 colonies of each vaccine onto BHI plates supplemented with streptomycin plus erythromycin (marker for antigen expression cassette integrated at the *tRNA<sup>arg</sup>* locus) or streptomycin plus chloramphenicol (marker for antigen expression cassette integrated at the *comK* locus). To assay vaccine stability in macrophage-like cells, we infected monolayers of J774A.1 cells with rLm vaccines in the absence of antibiotic selection for 5.5 hours; lysed the cells; serially diluted the lysates and plated

them on BHI agar supplemented with various antibiotics; cultured the plates at 37°C for 2 days; and counted the colonies.

### Immunization of mice and intracellular cytokine staining of mouse spleen and lung cells.

To determine the immunogenicity of rLm5Ag as a standalone vaccine, we immunized C57BL/6 and BALB/c mice, 4/group, s.q. at Weeks 0, 4, and 8 with  $2 \times 10^6$  Colony Forming Units (CFU) of the Lm vector (LmVector) or rLm5Ag (expressing the fusion protein of ActAN-Mpt64-EsxH-EsxA-EsxB-r30 from the *tRNA<sup>arg</sup>* locus); euthanized the mice at 6 days post the last immunization; prepared single cell suspensions of spleen and lung cells; stimulated the single cell suspensions with various Mtb antigens for 6 h or 22 h; and assayed T-cell immunity by intracellular cytokine staining (ICS) using an eight-color flow cytometry panel to analyze simultaneously multiple cytokines at the single-cell level as described by us previously <sup>22,23</sup>.

### Immunization and aerosol challenge of mice with virulent Mtb Erdman strain.

Groups of BALB/c or C57BL/6 mice, 8/group, were vaccinated intradermally (i.d.), intramuscularly (i.m.), intranasally (i.n.), or s.q. two or three times, 3 or 4 weeks apart, with  $10^6$  CFU of LmVector or of multi-antigenic rLm vaccines; challenged 3 or 4 weeks later by exposure to an aerosol generated by a nebulizer from a 10-ml single-cell suspension of Mtb Erdman strain ( $2.4 \times 10^4$  CFU/ml) for 30 min followed by settling for 5 min; euthanized at 10 weeks post challenge; and spleens and right lungs removed and

assayed for bacillus burden as described by us previously <sup>22</sup>. Control mice were sham vaccinated i.d. with PBS or immunized i.d. or i.n. with  $1 \times 10^6$  CFU BCG at Week 0.

### Immunization of guinea pigs and lymphocyte proliferation assay of their spleen and lung cells

Guinea pigs (male Hartley), 4/group, were immunized i.d. at Weeks 0, 3, and 6 with  $10^6$  CFU of the LmVector or rLm9Ag, bled and euthanized 6 days post the last immunization. Spleens and lungs were removed and single cell suspensions of spleen and lung cells prepared and stimulated with or without Mtb antigens. Lymphocyte proliferation using Flow cytometry analysis was assayed as described below.

Briefly, single cell suspensions of  $1 \times 10^7$  spleen and lung cells were stained with  $1 \mu\text{M}$  Cell Tracer Violet (CTV, ThermoFisher, labeling cells to trace multiple generations using dye dilution by flow cytometry) for 10 min at  $37^\circ\text{C}$  and washed with Phosphate buffer saline (PBS) supplemented with 5% fetal bovine serum. CTV treated cells were resuspended in T cell medium <sup>22</sup>, adjusted to  $5 \times 10^7$  cells/ml, seeded in 96-well round-bottom plates (NUNC) ( $5 \times 10^6$  cells per 0.1 ml per well) and incubated with or without 15-mer peptide pools of each of the 9 Mtb antigens (Ag85B, EspA, EsxA, EsxB, EsxH, EsxN, Mpt64, PPE68, TB8.4) ( $1 \mu\text{g/ml}$  per peptide) (PepMix, JPT Peptide Technologies, Berlin, Germany) or PPD ( $5 \mu\text{g/ml}$ ). Cells incubated with T-cell medium alone served as a negative control and cells incubated with ConA ( $5 \mu\text{g/ml}$ ) served as a positive control. After 4-days incubation at  $37^\circ\text{C}$  in a  $\text{CO}_2$  incubator, cells were collected, washed with PBS, and stained with LIVE/DEAD Fixable Near-IR Dead Cell (LD-NIR) (ThermoFisher),



followed by staining with cell surface markers of PanT-APC (BioRad), CD4-PE (BioRad), and CD8-FITC (BioRad). Lymphocyte proliferation was analyzed as loss of CTV staining (CTV<sup>Low</sup>) using an AttuneNxt flow cytometer (ThermoFisher). Data were analyzed using FlowJo software. Initial gating of the events included lymphocytes based on forward scatter vs. side scatter pattern, followed by selection for singlet cells, live PanT<sup>+</sup> cells, and subsequently CD4<sup>+</sup> and CD8<sup>+</sup> T cells. Proliferating live CD4<sup>+</sup> and CD8<sup>+</sup> T cells were identified by loss of CTV staining (CTV<sup>Low</sup>) on each of the cell populations. The gates for each cell population were determined by using the cells incubated without addition of antigen and verified by cells incubated with addition of ConA.

### Immunization and aerosol challenge of guinea pigs with virulent Mtb Erdman strain.

Groups of guinea pigs (Hartley, male), 10/group, were vaccinated i.d. once with  $10^3$  CFU BCG (positive control), or s.q. three times at Weeks 0, 3, and 6 with one dose ( $10^6$ ) of LmVector (negative control), or 3 escalating doses ( $10^5$ ,  $10^6$ , and  $10^7$ ) of rLm9Ag; challenged at Week 10 by aerosol with Mtb Erdman strain ( $2.4 \times 10^4$  CFU/ml); euthanized at Week 20; and spleens and right lungs removed and assayed for bacillus burden as described by us previously <sup>22</sup>.

## Vaccine stability after passaging in guinea pigs

Guinea pigs were immunized subcutaneously at the back of the neck area with  $1 \times 10^6$  rLm9Ag vaccine diluted in 0.1 ml PBS. At 0, 1, 2, 4, and 8 days post immunization, 2 guinea pigs were euthanized at each time point; the skin at the immunization site ( $\sim 1 \text{ cm}^2$ ), spleen, lung and liver of each animal were removed and homogenized in PBS; and the homogenates were serially diluted in PBS and plated onto BHI agar plates supplemented with streptomycin (200  $\mu\text{g/ml}$ ). The plates were incubated for 2 days at  $37^\circ\text{C}$  in a  $\text{CO}_2$  incubator. Bacterial colonies were recovered from plates of the various tissue homogenates at 0, 1, 2, and 4 days, but not at 8 days post immunization.

Recovered colonies were randomly selected and inoculated into 1 ml BHI broth plus streptomycin (200  $\mu\text{g/ml}$ ) and grown overnight without agitation. The bacteria were then collected from the overnight culture, lysed in SDS buffer, and subjected to SDS-PAGE and western blotting using a rabbit polyclonal antibody to Lm ActA (AK18).

## Statistics and reproducibility

Two-way ANOVA with Sidak's multiple comparisons test was performed to determine significance in comparisons of mean frequencies of cytokine-producing  $\text{CD4}^+$  and  $\text{CD8}^+$  T cells (Figs. 2, 3, 4 and 5); one-way ANOVA with Tukey's multiple comparisons test was used to determine mean organ CFU among mice in vaccinated and control groups (Figs. 6b, 7, 8). Two-way ANOVA with Sidak's multiple comparisons test was performed to determine significance in comparisons of mean frequencies of lymphocytes,  $\text{CD4}^+$ , and  $\text{CD8}^+$  T cells between guinea pigs vaccinated with LmVector

and those vaccinated with rLm9Ag (Fig. 9). Two-way ANOVA with Tukey's multiple comparisons test was used to determine significance in comparisons of percent weight change between guinea pigs vaccinated with LmVector and those vaccinated with rLm9Ag (Fig. 10b) and one-way ANOVA with Tukey's multiple comparisons test was used to determine significance in comparison of mean organ CFU among guinea pigs in vaccinated and control groups (Fig. 10c). All the statistical analyses were performed using Prism (9.2.0) software (GraphPad, San Diego, CA) except the Shapiro-Wilks tests. The Shapiro-Wilks test confirmed that all of the log scale CFU data in efficacy studies shown in Fig. 7, 8 and 10 have a normal distribution. Immunogenicity and efficacy studies were carried out in three animal models. The guinea pig immunogenicity study was repeated once.

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**Data availability:** All data supporting the findings of this study are available within the article and its supplementary information files or from the corresponding author upon reasonable request.

**Competing interests:** The authors declare no competing interests except M.A.H. and Q.J. are inventors on a patent application and on U.S. Patent 10,010,595 filed by UCLA on *Listeria monocytogenes*-vectored TB vaccines described in this and previous studies.

**Author contributions:** M.A.H. and Q.J. conceived the project; M.A.H. oversaw the project and obtained funding; Q.J., S.M.G, and S.N. conducted immunology and efficacy studies in mice and in guinea pigs; Q.J. processed data; Q.J. and M.A.H. wrote and revised the manuscript. All authors reviewed the manuscript in final form.

## Figure Legends

**Fig. 1. Construction and verification of rLm9Ag vaccine.** **a.** Diagram of rLm5Ag and rLm9Ag vaccine candidates. Expression cassettes ActAN-5Ag for rLm5Ag (top panel) and ActAN-5AgII for rLm5AgII (middle panel) driven by the Lm *actA* promoter were integrated at the *tRNA<sup>arg</sup>* loci of the corresponding rLm chromosome; expression cassettes ActAN-5Ag and ActAN-5AgII for rLm9Ag were integrated at the *comK* and *tRNA<sup>arg</sup>* loci of the Lm chromosome, respectively (bottom panel). **b.** Expression of heterologous and homologous proteins by rLm vaccine candidates grown in broth medium. Stocks of rLm vaccine candidates were inoculated into Brain Heart Infusion (BHI) medium and grown overnight at 37°C with agitation. The overnight culture was collected by centrifugation, resuspended in PBS supplemented with Halt Proteinase Inhibitor (Thermo Fisher Scientific), and lysed in SDS buffer. Equivalent amounts of bacterial lysates of each rLm were analyzed by Western blotting sequentially using rabbit polyclonal antibody to ActAN (pAb AK18) (top panel), rabbit polyclonal antibody to Mtb EsxH (middle panel); and monoclonal antibody to Lm P60 (serving as a loading control). The membrane was stripped before re-probing. **c.** The murine macrophage-like (J774A.1) cells were uninfected (UI) or infected at a MOI of 10 with LmVector or an rLm vaccine expressing Mtb 5Ag, 5AgII, or 9Ag that had been grown to stationary phase. At 5.5 h post-infection, the infected cells were lysed and subjected to Western blotting using polyclonal antibody AK18 (upper panel); the membrane was stripped and re-probed with a monoclonal antibody to Lm P60 and a monoclonal antibody to cellular protein  $\beta$ -actin (lower panel), as indicated on the right border of each panel. In both panels b and c: M, protein standards; Lane 1, LmVector (LmV); Lanes 2 and 3, two

clones of rLm5Ag expressing ActAN-Mpt64-EsxH-EsxA-EsxB-r30 from the *tRNA<sup>arg</sup>* locus; Lane 4, rLm5AgII expressing ActAN-Mpt64-EsxN-PPE68-EspA-TB8.4 from the *tRNA<sup>arg</sup>* locus; Lanes 5 and 6, two clones of rLm9Ag expressing a total of 9 different antigens, ActAN-5Ag from the *comK* locus and ActAN-5AgII from the *tRNA<sup>arg</sup>* locus; lane 7 (Panel c only), uninfected cells. On the left border of each panel are listed the sizes (kDa) of the molecular mass standards. Proteins of interests are indicated with arrows to the right of the protein band. Estimated Mw of ActAN-5AgII: 118-kDa; ActAN-5Ag: 94-kDa; Lm P60: 60 kDa; and  $\beta$ -actin, 42 kDa. The full-length images for b and c are shown in Supplementary Figure 1.

**Fig. 2. Frequency of cytokine-expressing CD4<sup>+</sup> T cells in the lungs and spleens of C57BL/6 mice immunized three times with LmVector or rLm5Ag.** C57BL/6 mice (n = 4/group) were immunized s.q. three times with LmVector (black) or rLm5Ag (pink) at Weeks 0, 4, and 8. Six days after the last immunization, mice were euthanized; their lungs and spleens removed; single cell suspensions prepared and stimulated with individual proteins of 23.5/Mpt64 (a, i), TB10.4/EsxH (b, j), ESAT6/EsxA (c, k), CFP10/EsxB (d, l), r30/Ag85B (e, m), or pool of 5 antigens (5Ag pool) (f, n), or PPD (g, o) in the presence of anti-CD28 monoclonal antibody for 6 h; GolgiPlug (protein transport inhibitor containing Brefeldin A) diluted in T-cell medium was added to all wells; additional cells incubated with PMA and Golgiplug for 4 hours served as positive control (h, p). The cells were assayed by excluding dead cells followed by staining for surface markers of CD4 and CD8 followed by CD3 and intracellular markers of IFN- $\gamma$ , TNF- $\alpha$ , IL-2, and IL-17A, as indicated below each panel. Frequencies of live CD4<sup>+</sup> T

cells expressing any of the four cytokines were analyzed by FlowJo 10 software. Values are the mean  $\pm$  standard error of the mean (SEM). \*,  $P < 0.05$ ; and \*\*,  $P < 0.01$  by two-way ANOVA with Sidak's post multiple comparisons test. A similar experiment was repeated in C57BL/6 mice.

**Fig. 3. Frequency of polyfunctional cytokine-expressing CD4<sup>+</sup> T cells in the lungs and spleens of C57BL/6 mice immunized three times with LmVector or rLm5Ag.**

C57BL/6 mice ( $n = 4$ /group, the same mice as shown in Fig. 2) were immunized three times s.q. with LmVector (black) or rLm5Ag (pink) at Weeks 0, 4, and 8. One week after the last immunization, animals were euthanized; their lung and spleen cells processed as described in the legend to Fig. 2. The lung and spleen cells were stimulated with r30/Ag85B (a, e), pool of 5 antigens (5Ag pool) (b, f), PPD (c, g), or PMA (positive control, (d, h), and assayed by intracellular cytokine staining for surface markers of CD4 and CD8 followed by CD3 and intracellular markers of IFN- $\gamma$ , TNF- $\alpha$ , IL-2, and IL-17A. The frequencies of live CD4<sup>+</sup> T cells producing any of the 15 possible combinations of the four cytokines (IFN- $\gamma$ , TNF- $\alpha$ , IL-2, and IL-17A) were uniquely distinguished by using Boolean gates of FlowJo 10 software, as indicated below each panel. Values are the mean  $\pm$  SEM. \*\*,  $P < 0.01$ ; \*\*\*,  $P < 0.001$ ; and \*\*\*\*,  $P < 0.0001$  by two-way ANOVA with Sidak's post multiple comparisons test.

**Fig. 4. Frequency of cytokine-expressing CD8<sup>+</sup> T cells in the lungs and spleens of C57BL/6 mice immunized three times with LmVector or rLm5Ag.**

C57BL/6 mice ( $n = 4$ /group, the same mice as shown in Fig. 2) were immunized three times s.q. with

LmVector (black) or rLm5Ag (pink) at Weeks 0, 4, and 8. One week after the last immunization, animals were euthanized and their lung and spleen cells processed as described in the legend to Fig. 2. The lung and spleen cells were stimulated with individual proteins of 23.5/Mpt64 (a, i), TB10.4/EsxH (b, j), ESAT6/EsxA (c, k), CFP10/EsxB (d, l), or r30/Ag85B (e, m), or pool of 5 antigens (5Ag pool) (f, n), PPD (g, o), or PMA (h, p), and assayed by intracellular cytokine staining for surface markers of CD4 and CD8 followed by CD3 and intracellular markers of IFN- $\gamma$ , TNF- $\alpha$ , IL-2, and IL-17A. Frequencies of CD8<sup>+</sup> T cells expressing any of the four cytokines are analyzed by FlowJo 10 software. Values are the mean  $\pm$  SEM. \*,  $P < 0.05$ ; \*\*,  $P < 0.01$ ; \*\*\*,  $P < 0.001$  by two-way ANOVA with Sidak's post multiple comparisons test.

**Fig. 5. Frequency of polyfunctional cytokine-expressing CD8<sup>+</sup> T cells in the lungs of C57BL/6 mice immunized three times with LmVector or rLm5Ag.** C57BL/6 mice (n = 4/group, the same mice as shown in Fig. 2) were immunized three times s.q. with LmVector (black) or rLm5Ag (pink) at Weeks 0, 4, and 8. One week after the last immunization, animals were euthanized and their lung cells processed as described in the legend to Fig. 2. The lung cells were stimulated with individual proteins of 23.5/Mpt64 (a) or TB10.4/EsxH (b), or pool of 5 antigens (5Ag pool) (c) or PPD (d), and assayed by intracellular cytokine staining for surface markers of CD4 and CD8 followed by CD3 and intracellular markers of IFN- $\gamma$ , TNF- $\alpha$ , IL-2, and IL-17A. The frequencies of live CD8<sup>+</sup> T cells producing any of the 15 possible combinations of the four cytokines (IFN- $\gamma$ , TNF- $\alpha$ , IL-2, and IL-17A) were uniquely distinguished by using Boolean gates of



FlowJo 10 software, as indicated below each panel. Values are the mean  $\pm$  SEM. \*\*\*\*,  $P < 0.0001$  by two-way ANOVA with Sidak's post multiple comparisons test.

**Fig. 6. Efficacy of immunization with rLm vaccines expressing 5 Mtb antigens**

**against Mtb aerosol challenge. a. Top Panel:** C57BL/6 mice ( $n = 8/\text{group}$ ) were immunized i.d. with PBS (Sham), i.d. or i.n with BCG at Week 0, or immunized i.d. or i.n. twice at Weeks 7 and 10 with rLm5Ag\* (rLm30 + rLm4Ag), challenged at Week 13 with aerosolized Mtb Erdman strain (average of 24 CFU delivered to the lung of each animal, as assayed at Day 1 post challenge) and euthanized at Week 23. **Middle and Bottom Panels:** Lungs and spleens of mice immunized i.d. (left panels) or i.n. (right panels) were assayed for organ bacterial burden. Shown are means  $\pm$  SEM. Each symbol represents one mouse. \*,  $P < 0.05$ ; \*\*,  $P < 0.01$ ; \*\*\*,  $P < 0.001$  and \*\*\*\*,  $P < 0.0001$  by 2-tailed non-paired t-test (Prism). **b. Top Panel:** BALB/c mice (8/group) were immunized i.d. with PBS or i.d. with BCG at Week 0, or immunized i.m. twice (x2) at Weeks 14 and 18 or three times (x3) at Weeks 10, 14, and 18 with rLm5Ag\*. The mice were then challenged with aerosolized Mtb Erdman strain (average of 30 CFU delivered to lung of each animal, as assayed at Day 1 post challenge) at Week 22 and euthanized at Week 32. **Middle and Bottom Panels:** Lungs and spleens were assayed for organ bacterial burden. Shown are means  $\pm$  SEM. Each symbol represents one mouse. \*,  $P < 0.05$ ; \*\*,  $P < 0.01$ ; \*\*\*,  $P < 0.001$ ; and \*\*\*\*,  $P < 0.0001$  by one-way ANOVA with Tukey's multiple comparisons test (Prism 9.2.0).

**Fig. 7. Efficacy of immunization with rLm5Ag and rLm9Ag vaccines against Mtb**

**aerosol challenge in C57BL/6 mice. a.** C57BL/6 mice ( $n = 8/\text{group}$ ) were

unimmunized (UI), vaccinated i.d. with  $5 \times 10^5$  CFU BCG at Week 0, or vaccinated s.q. three times with LmVector, rLm30, rLm5Ag\* (rLm30 + rLm4Ag), rLm5Ag (a single vaccine expressing the same 5 Mtb antigens as rLm5Ag\*), or rLm9Ag (clones #1 and #3) at Weeks 0, 3, and 6. The mice were then challenged at Week 10 with aerosolized Mtb Erdman strain (average of 10 CFU delivered to the lungs of each animal, as assayed at Day 1 post challenge) and euthanized at Week 20. **b.** Afterwards, lungs (left) and spleens (right) were removed and assayed for bacillus burdens. Shown are means  $\pm$  SEM. Each symbol represents one mouse. \*,  $P < 0.05$ ; \*\*,  $P < 0.01$ ; and \*\*\*,  $P < 0.001$  by one-way ANOVA with Tukey's multiple comparisons test (Prism 9.2.0). A similar experiment was repeated in BALB/c mice.

**Fig. 8. Efficacy of immunization with rLm5Ag and rLm9Ag vaccines against Mtb aerosol challenge in BALB/c mice.** **a.** BALB/c mice ( $n = 8$ /group) were unimmunized (UI), vaccinated i.d. with  $5 \times 10^5$  CFU BCG at Week 0, or vaccinated s.q. three times with LmVector, rLm30, rLm5Ag\* (combination of rLm30 + rLm4Ag), rLm5Ag (a single vaccine expressing the same 5 Mtb antigens as rLm5Ag\*), or rLm9Ag (clones #1 and #3) at Weeks 0, 3, and 6. The mice were then challenged at Week 10 with aerosolized Mtb Erdman strain (average of 10 CFU delivered to the lungs of each animal, as assayed at Day 1 post challenge) and euthanized at Week 20. **b.** Afterwards, lungs (left) and spleens (right) were removed and assayed for bacillus burdens. Shown are means  $\pm$  SEM. Each symbol represents one mouse. \*,  $P < 0.05$ ; \*\*,  $P < 0.01$ ; \*\*\*,  $P < 0.001$ ; and \*\*\*\*,  $P < 0.0001$  by one-way ANOVA with Tukey's multiple comparisons test (Prism). A similar experiment was repeated in C57BL/6 mice.

**Fig. 9. Frequencies of all lymphocytes and proliferating CD4+ and CD8+ T cells in the lungs and spleens of guinea pigs immunized with LmVector or rLm9Ag.**

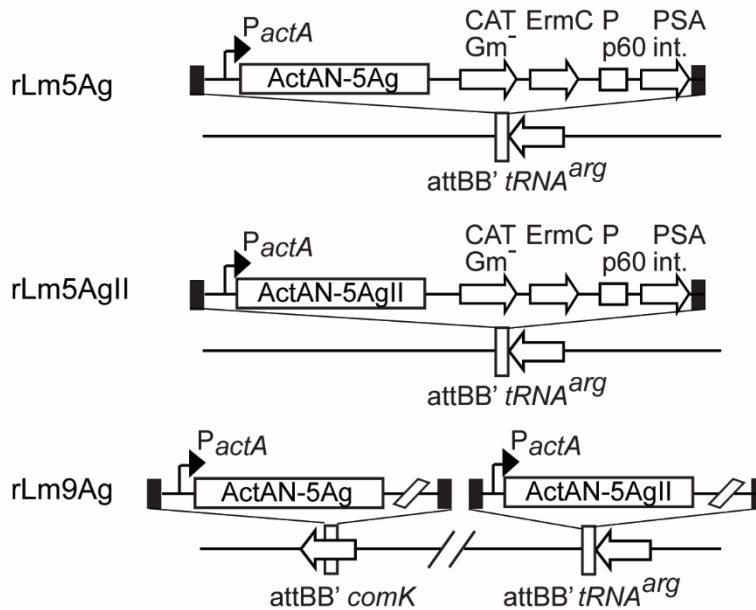
Guinea pigs (n = 4/group) were immunized three times s.q. with LmVector (black) or rLm9Ag (pink) at Weeks 0, 3, and 6. Six days after the last immunization, guinea pigs were euthanized; their spleens and lungs removed; single cell suspensions of spleen and lung cells prepared and treated with Cell Tracer Violet (CTV), followed by incubation with medium alone (negative control), medium with addition of 1 µg/ml per peptide of each of the 9 Mtb antigen peptide pools plus anti-CD28 monoclonal antibody (mAb), medium with addition of 5 µg/ml of PPD protein plus anti-CD28 mAb, or medium with addition of 5 µg/ml of ConA (positive control), as indicated beneath the horizontal axis, for 4 days. Subsequently, the cells were stained with Live/dead staining dye and surface markers of CD4-PE and CD8-FITC. Total lung lymphocytes (a) and total spleen lymphocytes (b) were gated by forward and side scatter patterns; proliferating lung CD4+ T cells (c), proliferating spleen CD4+ T cells (d), proliferating lung CD8+ T cells (e), and proliferating spleen CD8+ T cells (f) were gated by corresponding antibody staining. Values are means ± SEM. \*,  $P < 0.05$ ; \*\*,  $P < 0.01$ ; \*\*\*,  $P < 0.001$ ; \*\*\*\*,  $P < 0.0001$  by two-way ANOVA with Sidak's post multiple comparisons test (Prism 9.2.0). The experiment was repeated once with similar results.

**Fig. 10. Protective efficacy of rLm9Ag and rLm5Ag vaccines against Mtb aerosol challenge in guinea pigs.** a. Experimental design. Guinea pigs (n = 10/group) were immunized i.d. with  $10^3$  CFU BCG (Positive Control) at Week 0, or immunized s.q. three times with rLm9Ag at  $10^5$  (low dose, L),  $10^6$  (Medium dose, M), or  $10^7$  (High Dose, H)

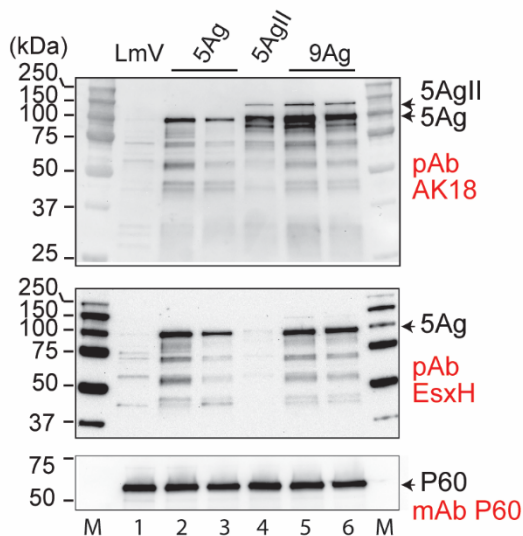
CFU, or with  $10^6$  rLm5Ag at Weeks 0, 3, and 6. The animals were then challenged at Week 10 with aerosolized Mtb Erdman strain. Afterwards, animals were monitored for signs of illness and weight change weekly for 10 weeks. At Week 10 post challenge, animals were euthanized and their spleen and right lungs were removed and assayed for bacillus burdens. **b.** Percent weight change post challenge. Shown are means  $\pm$  SEM. See Supplementary Figure 8 for data points. See Supplementary Table 2 for statistical analyses comparing LmVector, rLm5Ag, rLm9Ag, and BCG groups by two-way ANOVA with Tukey's multiple comparisons test (Prism 9.2.0). **c.** Right lung (left) and spleen bacterial burden. Shown are means  $\pm$  SEM. Each symbol represents one animal. \*,  $P < 0.05$ ; \*\*,  $P < 0.01$ ; \*\*\*,  $P < 0.001$ ; and \*\*\*\*,  $P < 0.0001$  by one-way ANOVA with Tukey's multiple comparisons test.

## Figures

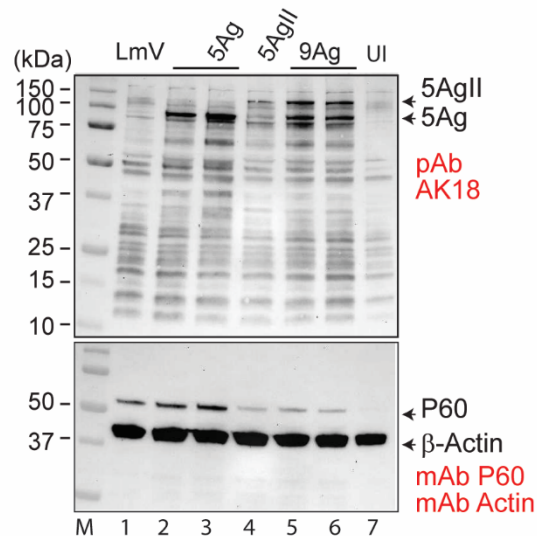
### a Diagram of rLm vaccine candidates



### b Protein expression in broth

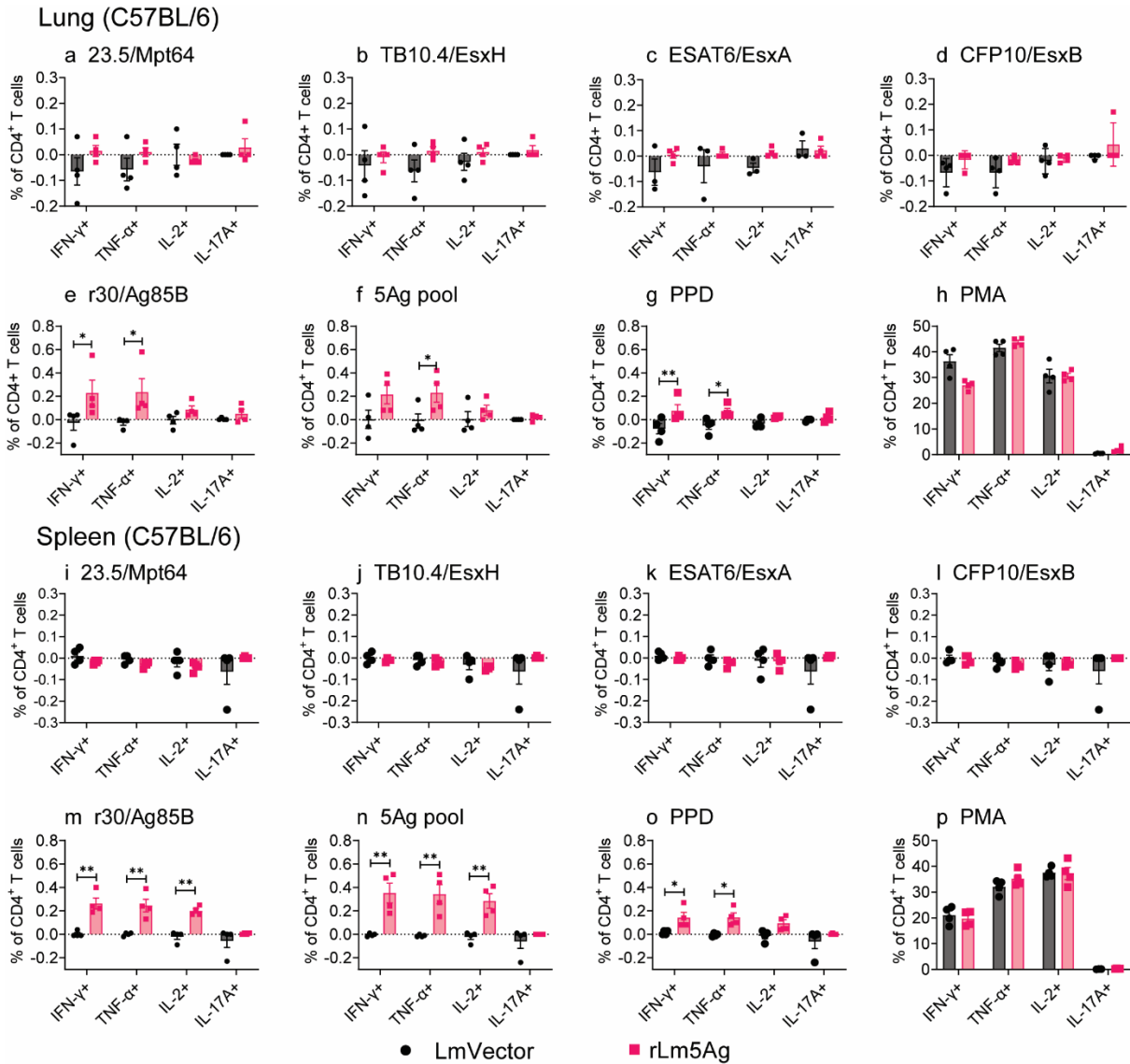


### c Protein expression in infected macrophages



**Fig. 1. Construction and verification of rLm9Ag vaccine.** **a.** Diagram of rLm5Ag and rLm9Ag vaccine candidates. Expression cassettes *ActAN-5Ag* for rLm5Ag (top panel) and *ActAN-5AgII* for rLm5AgII (middle panel) driven by the Lm *actA* promoter were integrated at the *tRNA<sup>arg</sup>* loci of the corresponding rLm chromosome; expression cassettes *ActAN-5Ag* and *ActAN-5AgII* for rLm9Ag were integrated at the *comK* and *tRNA<sup>arg</sup>* loci of the Lm chromosome, respectively (bottom panel). **b.** Expression of heterologous and homologous proteins by rLm vaccine candidates grown in broth

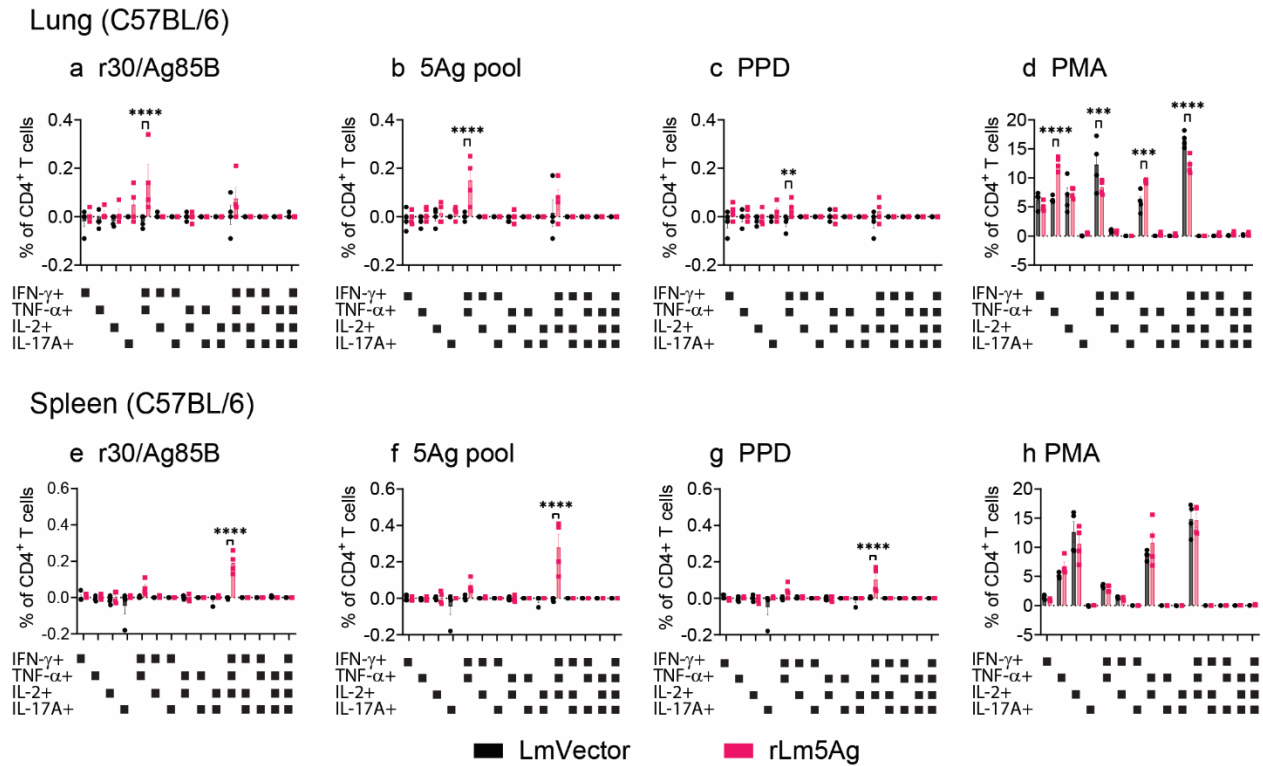
medium. Stocks of rLm vaccine candidates were inoculated into Brain Heart Infusion (BHI) medium and grown overnight at 37°C with agitation. The overnight culture was collected by centrifugation, resuspended in PBS supplemented with Halt Proteinase Inhibitor (Thermo Fisher Scientific), and lysed in SDS buffer. Equivalent amounts of bacterial lysates of each rLm were analyzed by Western blotting sequentially using rabbit polyclonal antibody to ActAN (pAb AK18) (top panel), rabbit polyclonal antibody to Mtb EsxH (middle panel); and monoclonal antibody to Lm P60 (serving as a loading control). The membrane was stripped before re-probing. **c.** The murine macrophage-like (J774A.1) cells were uninfected (UI) or infected at a MOI of 10 with LmVector or an rLm vaccine expressing Mtb 5Ag, 5AgII, or 9Ag that had been grown to stationary phase. At 5.5 h post-infection, the infected cells were lysed and subjected to Western blotting using polyclonal antibody AK18 (upper panel); the membrane was stripped and re-probed with a monoclonal antibody to Lm P60 and a monoclonal antibody to cellular protein  $\beta$ -actin (lower panel), as indicated on the right border of each panel. In both panels b and c: M, protein standards; Lane 1, LmVector (LmV); Lanes 2 and 3, two clones of rLm5Ag expressing ActAN-Mpt64-EsxH-EsxA-EsxB-r30 from the *tRNA<sup>arg</sup>* locus; Lane 4, rLm5AgII expressing ActAN-Mpt64-EsxN-PPE68-EspA-TB8.4 from the *tRNA<sup>arg</sup>* locus; Lanes 5 and 6, two clones of rLm9Ag expressing a total of 9 different antigens, ActAN-5Ag from the *comK* locus and ActAN-5AgII from the *tRNA<sup>arg</sup>* locus; lane 7 (Panel c only), uninfected cells. On the left border of each panel are listed the sizes (kDa) of the molecular mass standards. Proteins of interests are indicated with arrows to the right of the protein band. Estimated Mw of ActAN-5AgII: 118-kDa; ActAN-5Ag: 94-kDa; Lm P60: 60 kDa; and  $\beta$ -actin, 42 kDa. The full-length images for b and c are shown in Supplementary Figure 1.



**Fig. 2. Frequency of cytokine-expressing CD4<sup>+</sup> T cells in the lungs and spleens of C57BL/6 mice immunized three times with LmVector or rLm5Ag.** C57BL/6 mice (n = 4/group) were immunized s.q. three times with LmVector (black) or rLm5Ag (pink) at Weeks 0, 4, and 8. Six days after the last immunization, mice were euthanized; their lungs and spleens removed; single cell suspensions prepared and stimulated with individual proteins of 23.5/Mpt64 (a, i), TB10.4/EsxH (b, j), ESAT6/EsxA (c, k), CFP10/EsxB (d, l), r30/Ag85B (e, m), or pool of 5 antigens (5Ag pool) (f, n), or PPD (g, o) in the presence of anti-CD28 monoclonal antibody for 6 h; GolgiPlug (protein transport inhibitor containing Brefeldin A) diluted in T-cell medium was added to all wells; additional cells incubated with PMA and Golgiplug for 4 hours served as positive control (h, p). The cells were assayed by excluding dead cells followed by staining for surface markers of CD4 and CD8 followed by CD3 and intracellular markers of IFN- $\gamma$ ,

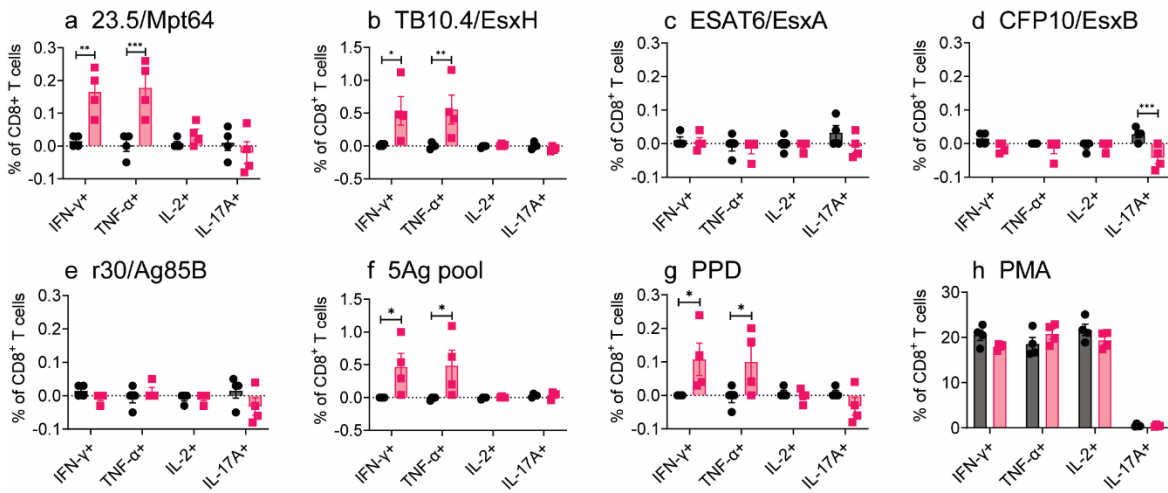
TNF- $\alpha$ , IL-2, and IL-17A, as indicated below each panel. Frequencies of live CD4+ T cells expressing any of the four cytokines were analyzed by FlowJo 10 software. Values are the mean  $\pm$  standard error of the mean (SEM). \*,  $P < 0.05$ ; and \*\*,  $P < 0.01$  by two-way ANOVA with Sidak's post multiple comparisons test. A similar experiment was repeated in BALB/c mice.



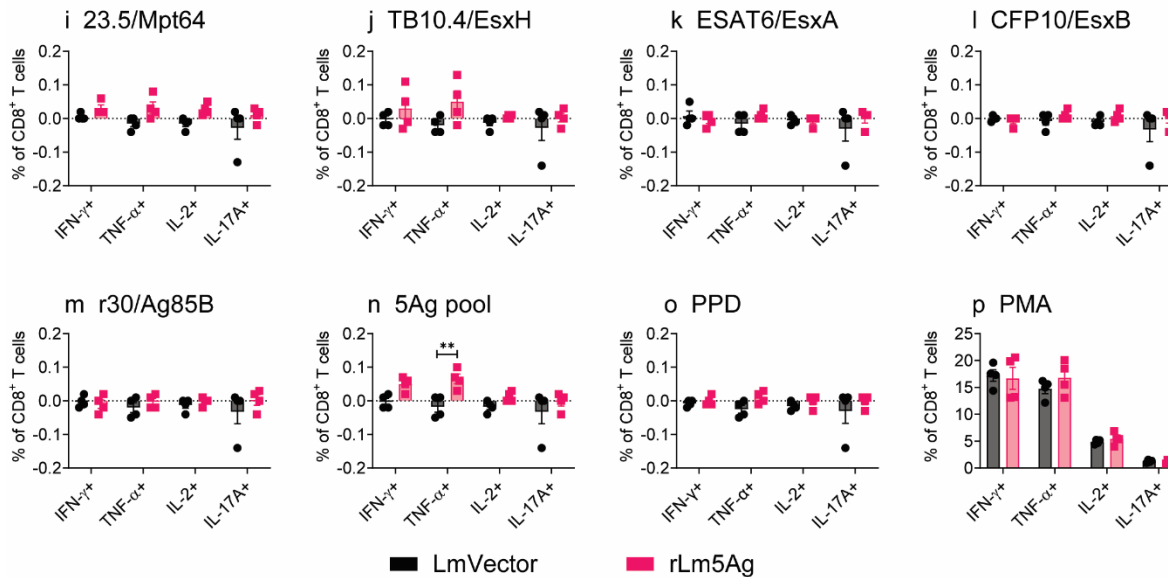


**Fig. 3. Frequency of polyfunctional cytokine-expressing CD4<sup>+</sup> T cells in the lungs and spleens of C57BL/6 mice immunized three times with LmVector or rLm5Ag.** C57BL/6 mice (n = 4/group, the same mice as shown in Fig. 2) were immunized three times s.q. with LmVector (black) or rLm5Ag (pink) at Weeks 0, 4, and 8. One week after the last immunization, animals were euthanized; their lung and spleen cells processed as described in the legend to Fig. 2. The lung and spleen cells were stimulated with individual proteins of r30/Ag85B (a, e), pool of 5 antigens (5Ag pool) (b, f), PPD (c, g) or PMA (positive control, d, h) and assayed by intracellular cytokine staining for surface markers of CD4 and CD8 followed by CD3 and intracellular markers of IFN- $\gamma$ , TNF- $\alpha$ , IL-2, and IL-17A. The frequencies of live CD4<sup>+</sup> T cells producing any of the 15 possible combinations of the four cytokines (IFN- $\gamma$ , TNF- $\alpha$ , IL-2, and IL-17A) were uniquely distinguished by using Boolean gates of FlowJo 10 software, as indicated below each panel. Values are the mean  $\pm$  SEM. \*\*,  $P < 0.01$ ; \*\*\*,  $P < 0.001$ ; and \*\*\*\*,  $P < 0.0001$  by two-way ANOVA with Sidak's post multiple comparisons test.

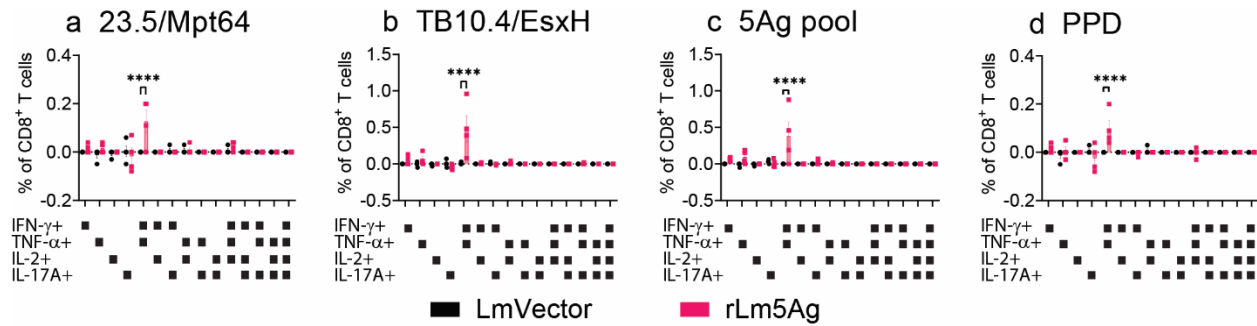
Lung (C57BL/6)



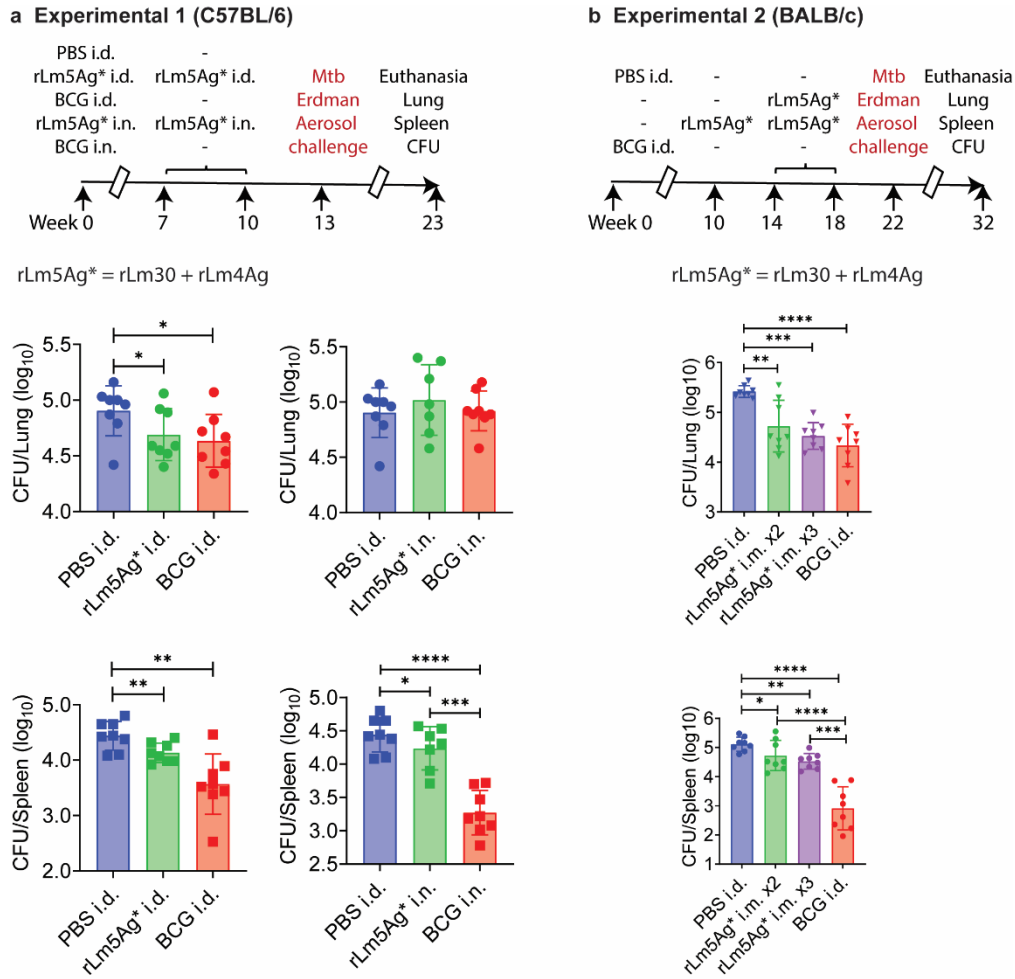
Spleen (C57BL/6)



**Fig. 4. Frequency of cytokine-expressing CD8<sup>+</sup> T cells in the lungs and spleens of C57BL/6 mice immunized three times with LmVector or rLm5Ag.** C57BL/6 mice (n = 4/group, the same mice as shown in Fig. 2) were immunized three times s.q. with LmVector (black) or rLm5Ag (pink) at Weeks 0, 4, and 8. One week after the last immunization, animals were euthanized and their lung and spleen cells processed as described in the legend to Fig. 2. The lung and spleen cells were stimulated with individual proteins of 23.5/Mpt64 (a, i), TB10.4/EsxH (b, j), ESAT6/EsxA (c, k), CFP10/EsxB (d, l), r30/Ag85B (e, m), or pool of 5 antigens (5Ag pool) (f, n) or PPD (g, o) or PMA (h, p), and assayed by intracellular cytokine staining for surface markers of CD4 and CD8 followed by CD3 and intracellular markers of IFN- $\gamma$ , TNF- $\alpha$ , IL-2, and IL-17A. Frequencies of CD8<sup>+</sup> T cells expressing any of the four cytokines are analyzed by FlowJo 10 software. Values are the mean  $\pm$  SEM. \*,  $P < 0.05$ ; \*\*,  $P < 0.01$ ; \*\*\*,  $P < 0.001$  by two-way ANOVA with Sidak's post multiple comparisons test.

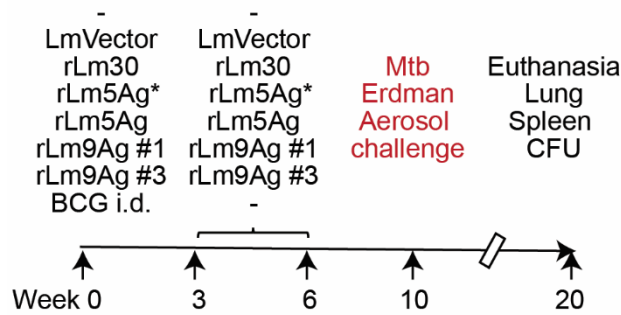


**Fig. 5. Frequency of polyfunctional cytokine-expressing CD8<sup>+</sup> T cells in the lungs of C57BL/6 mice immunized three times with LmVector or rLm5Ag.** C57BL/6 mice (n = 4/group, the same mice as shown in Fig. 2) were immunized three times s.q. with LmVector (black) or rLm5Ag (pink) at Weeks 0, 4, and 8. One week after the last immunization, animals were euthanized and their lung cells processed as described in the legend to Fig. 2. The lung cells were stimulated with individual proteins of 23.5/Mpt64 (a), TB10.4/EsxH (b), or pool of 5 antigens (5Ag pool) (c) or PPD (d) and assayed by intracellular cytokine staining for surface markers of CD4 and CD8 followed by CD3 and intracellular markers of IFN- $\gamma$ , TNF- $\alpha$ , IL-2, and IL-17A. The frequencies of live CD8<sup>+</sup> T cells producing any of the 15 possible combinations of the four cytokines (IFN- $\gamma$ , TNF- $\alpha$ , IL-2, and IL-17A) were uniquely distinguished by using Boolean gates of FlowJo 10 software, as indicated below each panel. Values are the mean  $\pm$  SEM. \*\*\*\*,  $P < 0.0001$  by two-way ANOVA with Sidak's post multiple comparisons test.

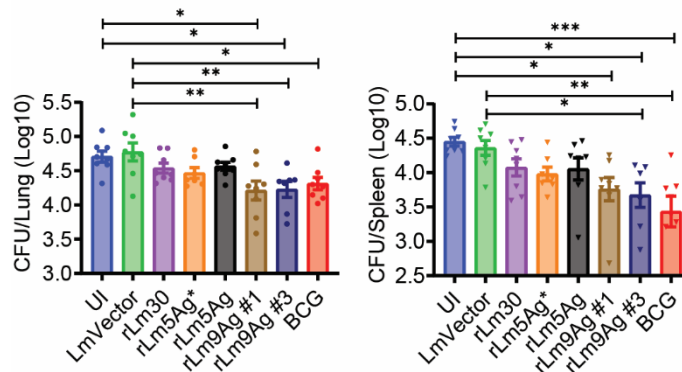


**Fig. 6. Efficacy of immunization with rLm vaccines expressing 5 Mtb antigens against Mtb aerosol challenge. a. Top Panel:** C57BL/6 mice ( $n = 8/\text{group}$ ) were immunized i.d. with PBS (Sham), i.d. or i.n. with BCG at Week 0, or immunized i.d. or i.n. twice at Weeks 7 and 10 with rLm5Ag\* (rLm30 + rLm4Ag), challenged at Week 13 with aerosolized Mtb Erdman strain (average of 24 CFU delivered to the lung of each animal, as assayed at Day 1 post challenge) and euthanized at Week 23. **Middle and Bottom Panels:** Lungs and spleens of mice immunized i.d. (left panels) or i.n. (right panels) were assayed for organ bacterial burden. Shown are means  $\pm$  SEM. Each symbol represents one mouse. \*,  $P < 0.05$ ; \*\*,  $P < 0.01$ ; \*\*\*,  $P < 0.001$  and \*\*\*\*,  $P < 0.0001$  by 2-tailed non-paired t-test (Prism). **b. Top Panel:** BALB/c mice (8/group) were immunized i.d. with PBS or i.d. with BCG at Week 0, or immunized i.m. twice (x2) at Weeks 14 and 18 or three times (x3) at Weeks 10, 14, and 18 with rLm5Ag\*. The mice were then challenged with aerosolized Mtb Erdman strain (average of 30 CFU delivered to lung of each animal, as assayed at Day 1 post challenge) at Week 22 and euthanized at Week 32. **Middle and Bottom Panels:** Lungs and spleens were assayed for organ bacterial burden. Shown are means  $\pm$  SEM. Each symbol represents one mouse. \*,  $P < 0.05$ ; \*\*,  $P < 0.01$ ; \*\*\*,  $P < 0.001$ ; and \*\*\*\*,  $P < 0.0001$  by one-way ANOVA with Tukey's multiple comparisons test (Prism 9.2.0).

### a Experimental design

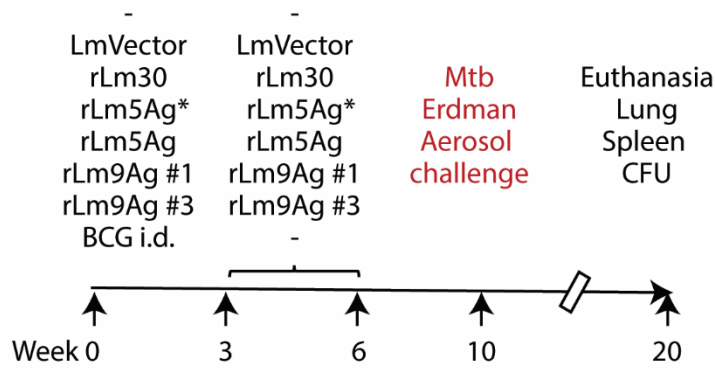


### b Organ CFUs

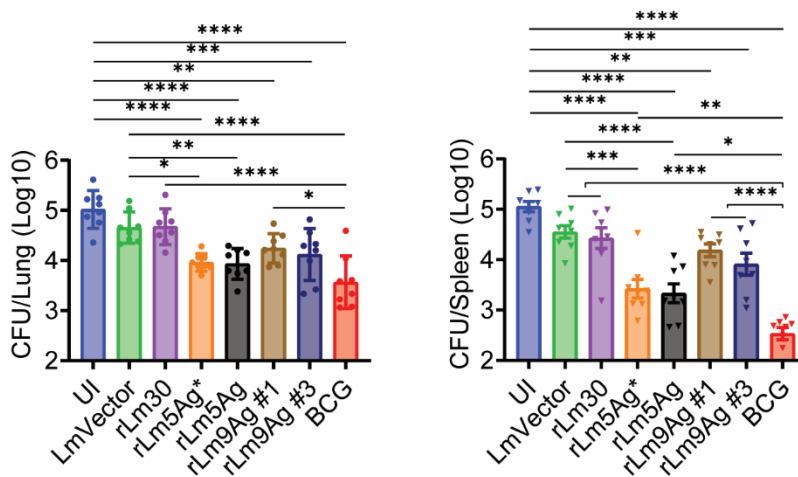


**Fig. 7. Efficacy of immunization with rLm5Ag and rLm9Ag vaccines against Mtb aerosol challenge in C57BL/6 mice.** **a.** C57BL/6 mice ( $n = 8/\text{group}$ ) were unimmunized (UI), vaccinated i.d. with  $5 \times 10^5$  CFU BCG at Week 0, or vaccinated s.q. three times with LmVector, rLm30, rLm5Ag\* (rLm30 + rLm4Ag), rLm5Ag (a single vaccine expressing the same 5 Mtb antigens as rLm5Ag\*), or rLm9Ag (clones #1 and #3) at Weeks 0, 3, and 6. The mice were then challenged at Week 10 with aerosolized Mtb Erdman strain (average of 10 CFU delivered to the lungs of each animal, as assayed at Day 1 post challenge) and euthanized at Week 20. **b.** Afterwards, lungs (left) and spleens (right) were removed and assayed for bacillus burdens. Shown are means  $\pm$  SEM. Each symbol represents one mouse. \*,  $P < 0.05$ ; \*\*,  $P < 0.01$ ; and \*\*\*,  $P < 0.001$  by one-way ANOVA with Tukey's multiple comparisons test (Prism 9.2.0). A similar experiment was repeated in BALB/c mice.

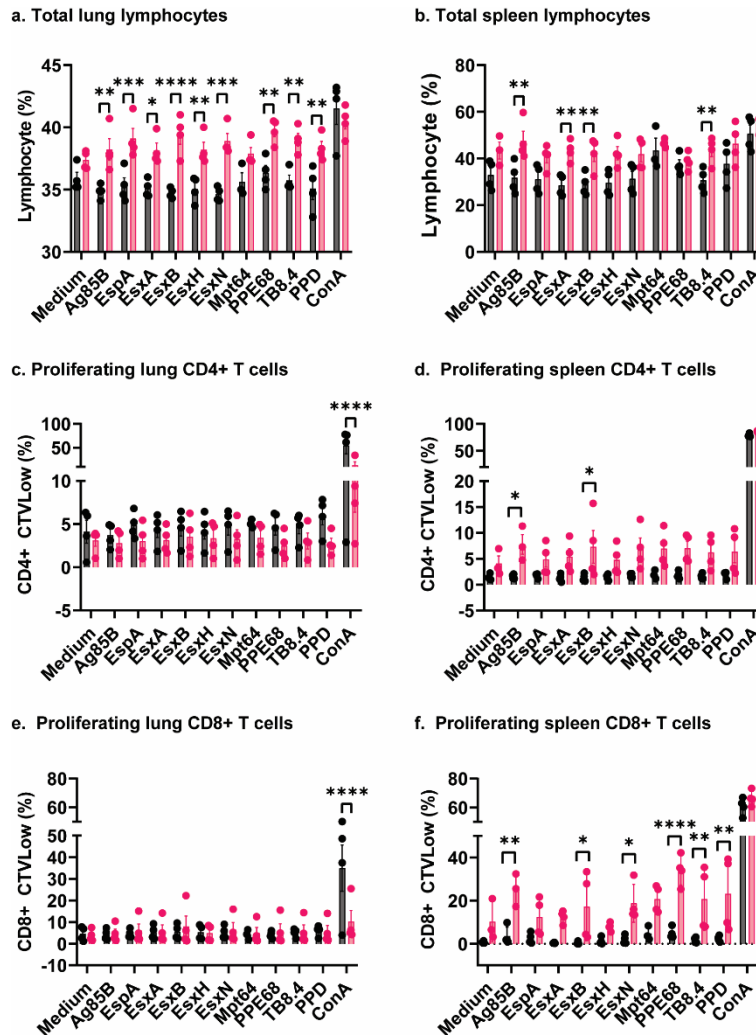
### a Experimental design



### b Organ CFUs



**Fig. 8. Efficacy of immunization with rLm5Ag and rLm9Ag vaccines against Mtb aerosol challenge in BALB/c mice.** **a.** BALB/c mice ( $n = 8/\text{group}$ ) were unimmunized (UI), vaccinated i.d. with  $5 \times 10^5$  CFU BCG at Week 0, or vaccinated s.q. three times with LmVector, rLm30, rLm5Ag\* (combination of rLm30 + rLm4Ag), rLm5Ag (a single vaccine expressing the same 5 Mtb antigens as rLm5Ag\*), or rLm9Ag (clones #1 and #3) at Weeks 0, 3, and 6. The mice were then challenged at Week 10 with aerosolized Mtb Erdman strain (average of 10 CFU delivered to the lungs of each animal, as assayed at Day 1 post challenge) and euthanized at Week 20. **b.** Afterwards, lungs (left) and spleens (right) were removed and assayed for bacillus burdens. Shown are means  $\pm$  SEM. Each symbol represents one mouse. \*,  $P < 0.05$ ; \*\*,  $P < 0.01$ ; \*\*\*,  $P < 0.001$ ; and \*\*\*\*,  $P < 0.0001$  by one-way ANOVA with Tukey's multiple comparisons test (Prism). A similar experiment was repeated in C57BL/6 mice.



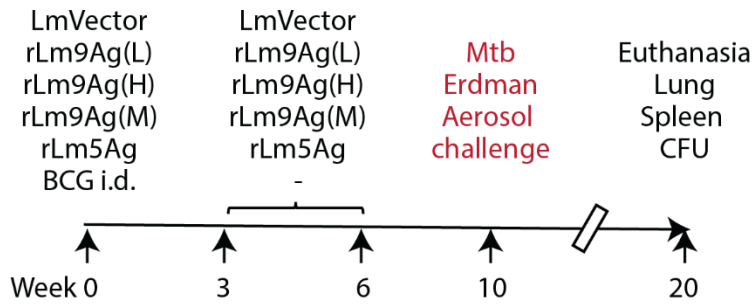
**Fig. 9. Frequencies of all lymphocytes and proliferating CD4+ and CD8+ T cells in the lungs and spleens of guinea pigs immunized with LmVector or rLm9Ag.**

Guinea pigs (n = 4/group) were immunized three times s.q. with LmVector (black) or rLm9Ag (pink) at Weeks 0, 3, and 6. Six days after the last immunization, guinea pigs were euthanized; their spleens and lungs removed; single cell suspensions of spleen and lung cells prepared and treated with Cell Tracer Violet (CTV), followed by incubation with medium alone (negative control), medium with addition of 1 µg/ml per peptide of each of the 9 Mtb antigen peptide pools plus anti-CD28 monoclonal antibody (mAb), medium with addition of 5 µg/ml of PPD protein plus anti-CD28 mAb, or medium with addition of 5 µg/ml of ConA (positive control), as indicated beneath the horizontal axis, for 4 days. Subsequently, the cells were stained with Live/dead staining dye and surface markers of CD4-PE and CD8-FITC. Total lung lymphocytes (a) and total spleen lymphocytes (b) were gated by forward and side scatter patterns; proliferating lung CD4+ T cells (c), proliferating spleen CD4+ T cells (d), proliferating lung CD8+ T cells (e), and proliferating spleen CD8+ T cells (f) were gated by corresponding antibody

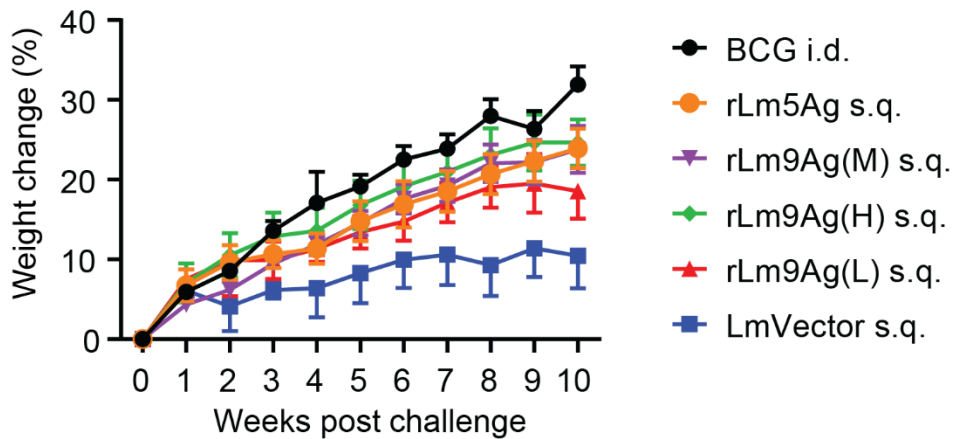
staining. Values are means  $\pm$  SEM. \*,  $P < 0.05$ ; \*\*,  $P < 0.01$ ; \*\*\*,  $P < 0.001$ ; \*\*\*\*,  $P < 0.0001$  by two-way ANOVA with Sidak's post multiple comparisons test (Prism 9.2.0).



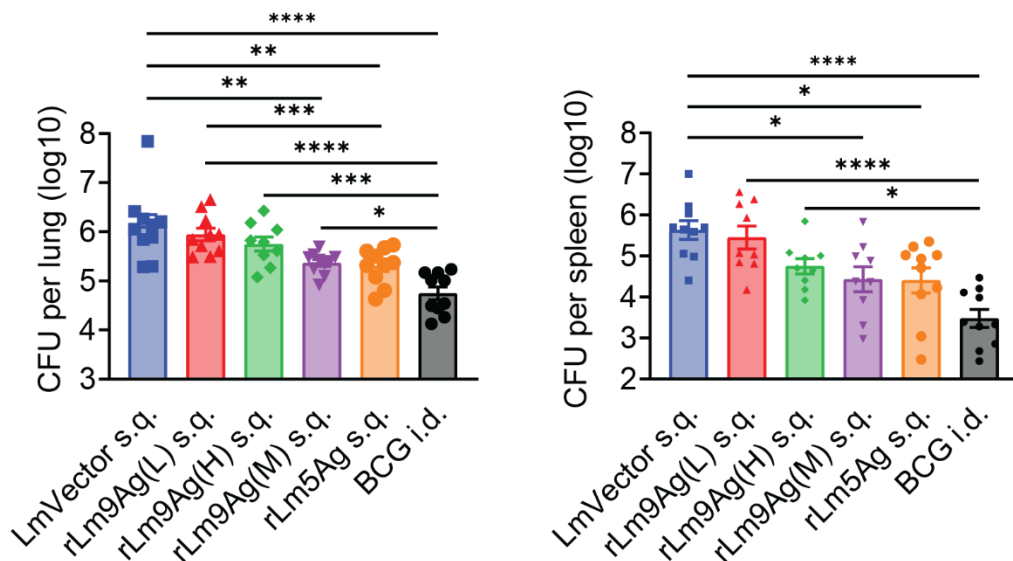
### a Experimental design



### b Percent weight change post challenge



### c Organ CFUs



**Fig. 10. Protective efficacy of rLm9Ag and rLm5Ag vaccines against Mtb aerosol challenge in guinea pigs.** a. Experimental design. Guinea pigs (n = 10/group) were immunized i.d. with  $10^3$  CFU BCG (Positive Control) at Week 0, or immunized s.q. three times with rLm9Ag at  $10^5$  (low dose, L),  $10^6$  (Medium dose, M), or  $10^7$  (High Dose, H)

CFU, or with  $10^6$  rLm5Ag at Weeks 0, 3, and 6. The animals were then challenged at Week 10 with aerosolized Mtb Erdman strain. Afterwards, animals were monitored for signs of illness and weight change weekly for 10 weeks. At Week 10 post challenge, animals were euthanized and their spleen and right lungs were removed and assayed for bacillus burdens. **b.** Percent weight change post challenge. Shown are means  $\pm$  SEM. See Supplementary Figure 8 for data points. See Supplementary Table 2 for statistical analyses comparing LmVector, rLm5Ag, rLm9Ag, and BCG groups by two-way ANOVA with Tukey's multiple comparisons test (Prism 9.2.0). **c.** Right lung (left) and spleen bacterial burden. Shown are means  $\pm$  SEM. Each symbol represents one animal. \*,  $P < 0.05$ ; \*\*,  $P < 0.01$ ; \*\*\*,  $P < 0.001$ ; and \*\*\*\*,  $P < 0.0001$  by one-way ANOVA with Tukey's multiple comparisons test.

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# Supplementary Materials

## **Listeria-vectored multi-antigenic tuberculosis vaccine protects C57BL/6 and BALB/c mice and guinea pigs against *Mycobacterium tuberculosis* challenge**

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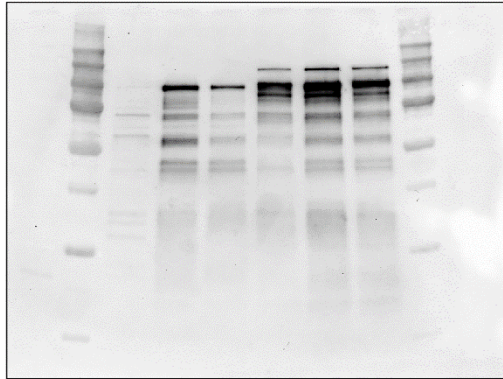
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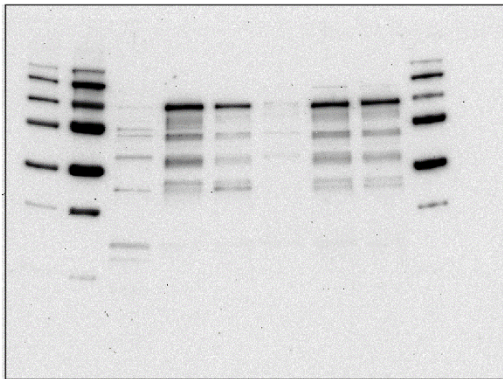
[MHorwitz@mednet.ucla.edu](mailto:MHorwitz@mednet.ucla.edu)

## Supplementary Figures

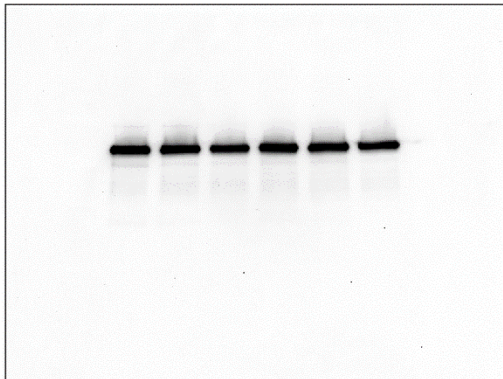
Full images for WB shown in Fig. 1b



PAb to  
AK18

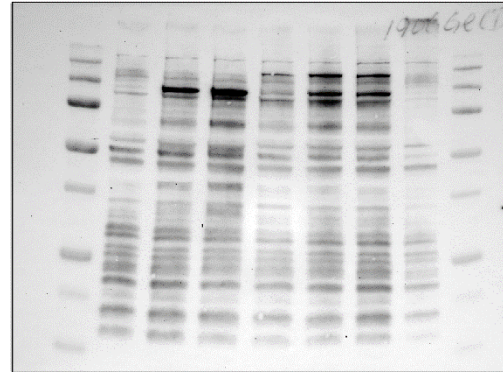


PAb to  
EsxH

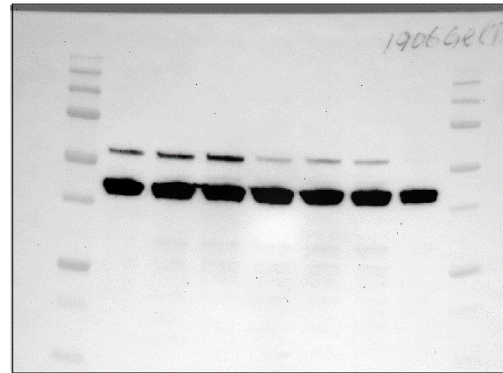


PAb to  
P60

Full image for WB shown in Fig. 1c

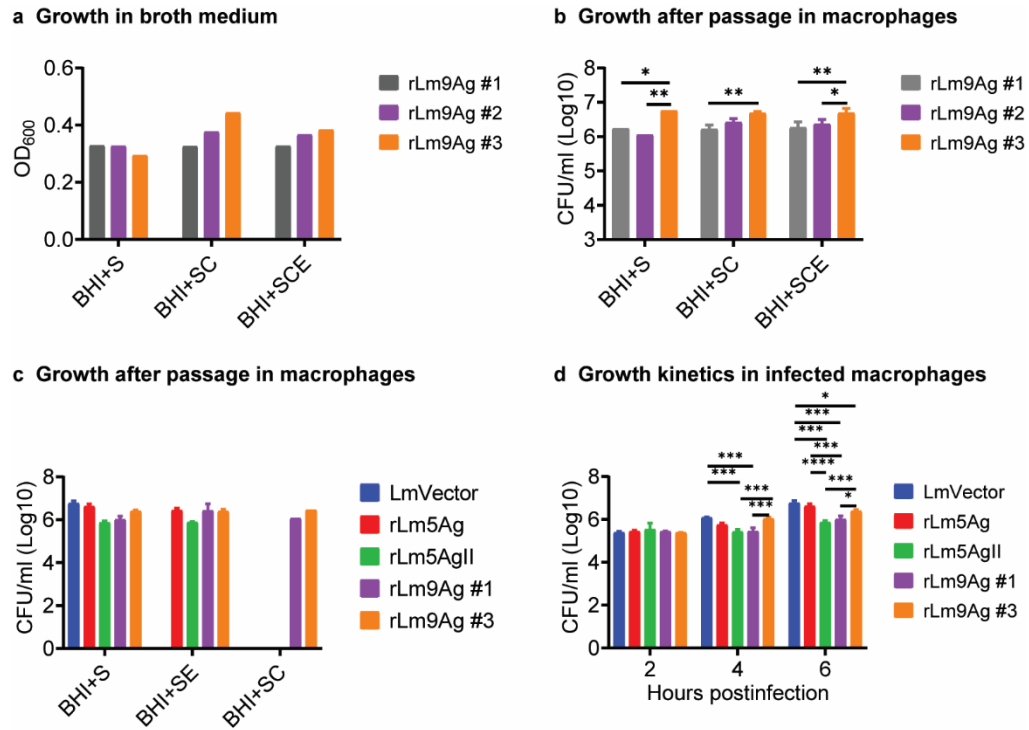


PAb to  
AK18

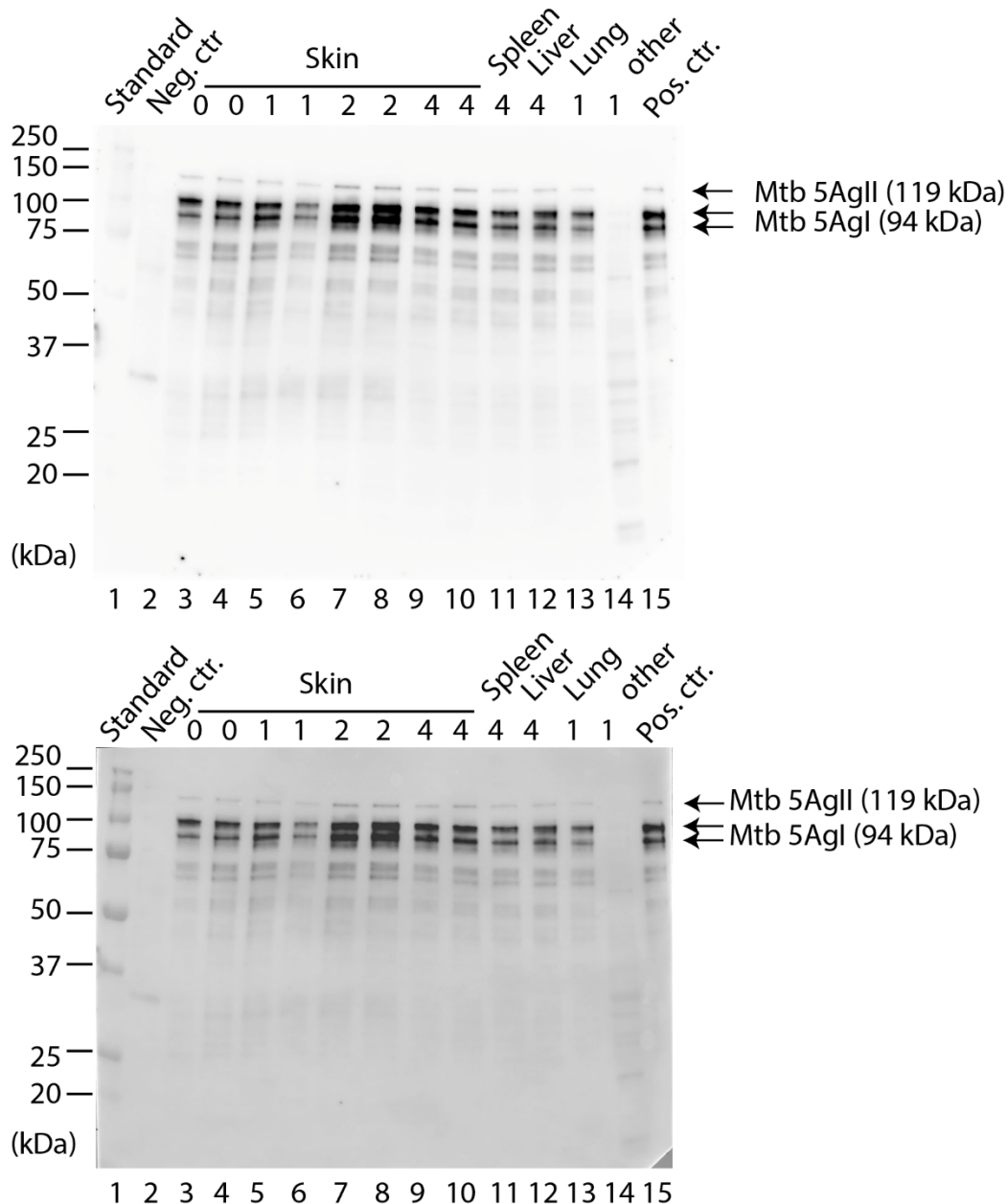


PAb to  
P60  
+ mAb to  
b-actin

**Supplementary Figure 1. Full images of Western blotting results shown in Figure 1b and 1c.**

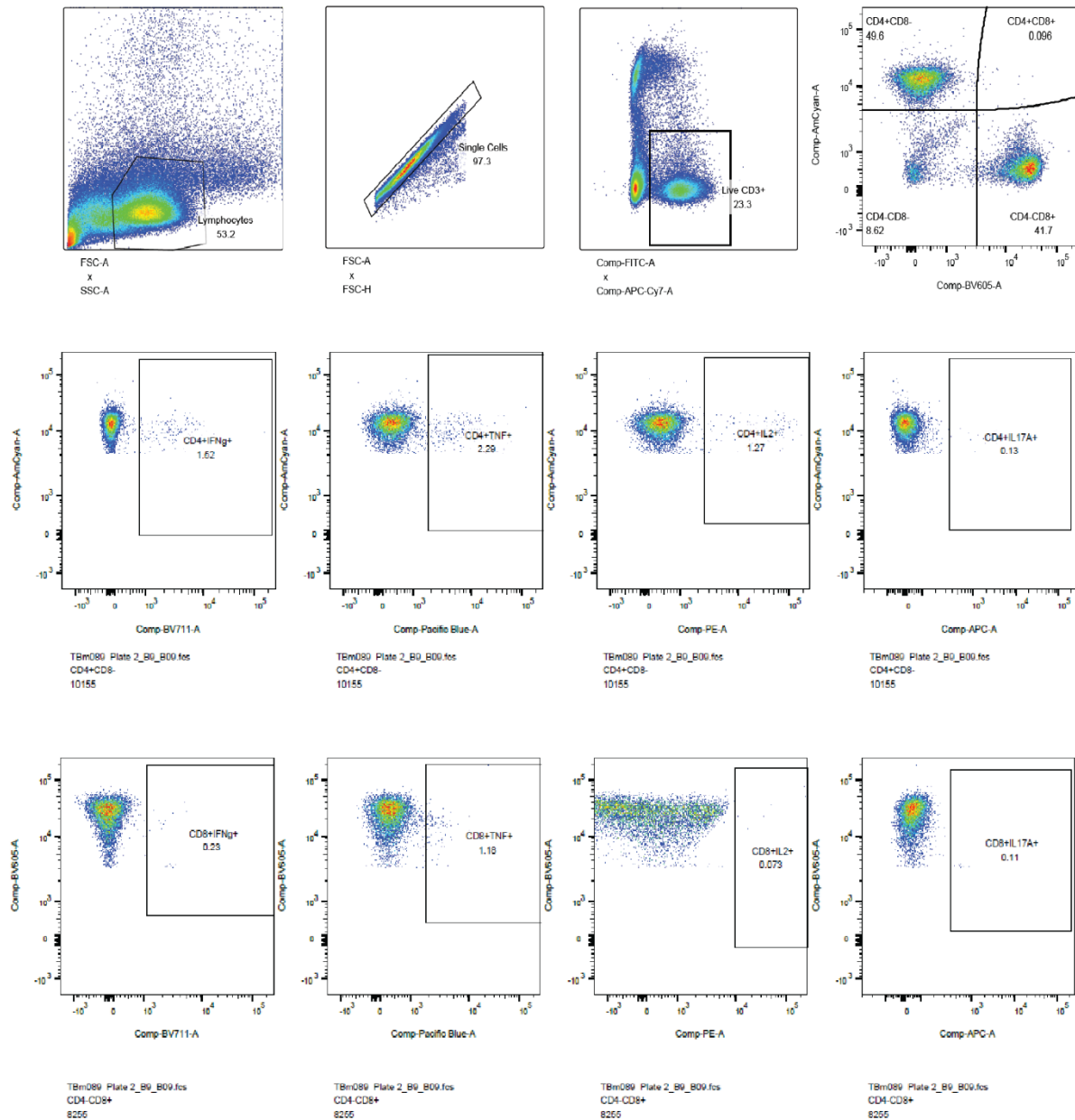


**Supplementary Figure 2. The antigen expression cassettes in the rLm chromosomes are stable in the presence or absence of antibiotic selection. a.** Each of the three clones of rLm9Ag were inoculated into Brain Heart Infusion (BHI) medium supplemented with Streptomycin (200  $\mu\text{g/ml}$ ) (BHI + S) (the Lm Vector is streptomycin resistant); Streptomycin (200  $\mu\text{g/ml}$ ) + Chloramphenicol (2.5  $\mu\text{g/ml}$ ) (BHI + SC); or Streptomycin (200  $\mu\text{g/ml}$ ) + Chloramphenicol (7.5  $\mu\text{g/ml}$ ) + Erythromycin (2.5  $\mu\text{g/ml}$ ) (BHI + SCE), as indicated below the horizontal axis. The cultures were grown overnight at 37°C with 5% CO<sub>2</sub> and optical density measured at 600 nm (OD<sub>600</sub>). **b & c.** Monolayers of J774A.1 cells were infected with the rLm indicated for 5.5 hours; the lysates serially diluted and plated on BHI agar supplemented with various antibiotics, as described in **a**; cultured at 37°C for 2 days; and the colonies (CFU) enumerated. **d.** Monolayers of J774A.1 cells were infected with LmVector or the rLm indicated with an MOI of 10. At 2, 4, and 6 h post infection, cells were lysed, the lysates plated on BHI agar plates without antibiotic selection, the plates incubated for 2 days, and CFU enumerated. Values are the mean  $\pm$  standard error of the mean (SEM). Values are mean log<sub>10</sub> CFU  $\pm$  SEM.



**Supplementary Figure 3. Protein expression by rLm9Ag after passage in guinea pigs.** Guinea pigs were immunized subcutaneously at the back of the neck area with  $1 \times 10^6$  rLm9Ag vaccine diluted in 0.1 ml PBS. At 0, 1, 2, 4, and 8 days post immunization, 2 guinea pigs were euthanized at each time point; the skin at the immunization site ( $1 \text{ cm}^2$ ), spleen, lung and liver of each animal were removed and homogenized in phosphate buffered saline; and the homogenates were serially diluted and plated onto BHI agar plates supplemented with streptomycin ( $200 \mu\text{g/ml}$ ) (BHI+Strep). The plates were incubated for 2 days at  $37^\circ\text{C}$  in a  $\text{CO}_2$  incubator. Bacterial colonies were recovered from plates of the various tissue homogenates at 0, 1, 2, and 4 days, but not at 8 days post immunization. Recovered colonies were randomly selected and inoculated into 1 ml BHI broth + strep and grown overnight without agitation. The

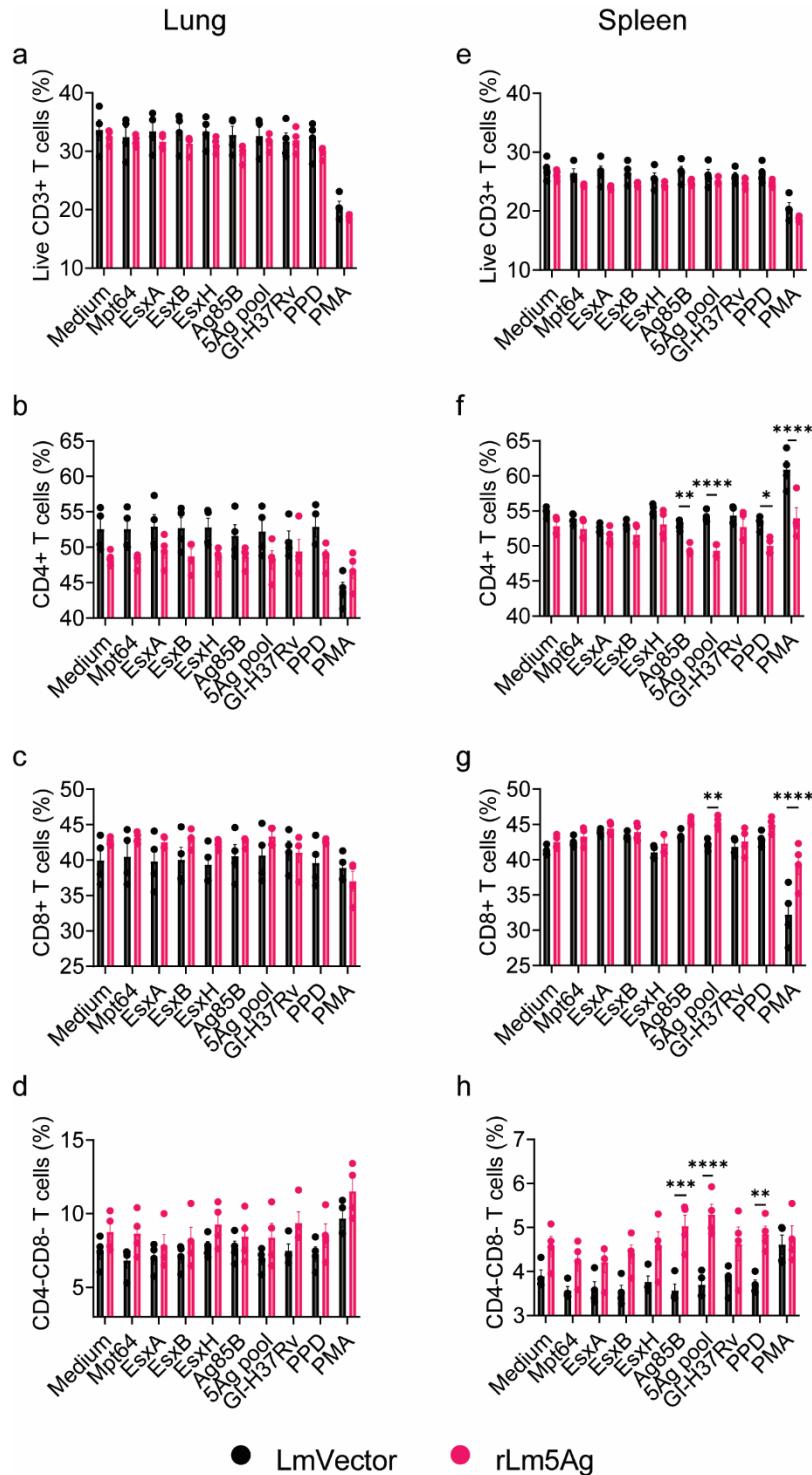
bacteria were then collected from the overnight culture, lysed in SDS buffer, and subjected to SDS-PAGE and western blotting using a rabbit polyclonal antibody to Lm ActA (AK18). Standard: Protein standards with their masses listed to the left of the panels. Neg. ctr: Lm vector negative control. Numbers (0, 1, 2, 4) on the top of the panels: Days post immunization. Other (lane 14): a contaminant from a BHI+strep plate, also serving as a negative control. Pos. ctr: rLm1939 vaccine stock positive control. Top panel, chemiluminescence. Bottom panel, composite of chemiluminescence and stain free gel, showing the protein standards. Arrows to the right of the panels indicate expected protein bands, Mtb 5Ag (ActAN-Mpt64-EsxH-EsxA-EsxB-r30, 94 kDa doublets) and 5AgII (ActAN-Mpt64-EsxN-PPE68-EspA-TB8.4, 119 kDa). Mtb 5Ag and Mtb5AgII share a common antigen of Mpt64.



**Supplementary Figure 4. Gating strategy of intracellular staining assay.** C57BL/6 mice ( $n = 4/\text{group}$ ) were immunized s.q. three times with LmVector or rLm5Ag at Weeks 0, 4, and 8. Six days after the last immunization, mice were euthanized; their spleens removed; single cell suspensions prepared and stimulated in T-cell medium without or with individual proteins 23.5/Mpt64, TB10.4/EsxH, ESAT6/EsxA, CFP10/EsxB, or r30/Ag85B; pool of these 5 antigens; or PPD in the presence of anti-CD28 monoclonal antibody for 2 h. Then, GolgiPlug (protein transport inhibitor containing Brefeldin A) diluted in T-cell medium was added to all wells and incubated with cells for an additional 4 h (total stimulation time of 6 h). Following *in vitro* stimulation, cells were harvested; washed with PBS; incubated with Live/Dead Fixable Near-IR Cell Stain (ThermoFisher) for 10 min at room temperature to identify dead cells; washed with Cell Staining Buffer (BioLegend) and then incubated with antibodies to CD4 (Clone RM4-5, conjugated with

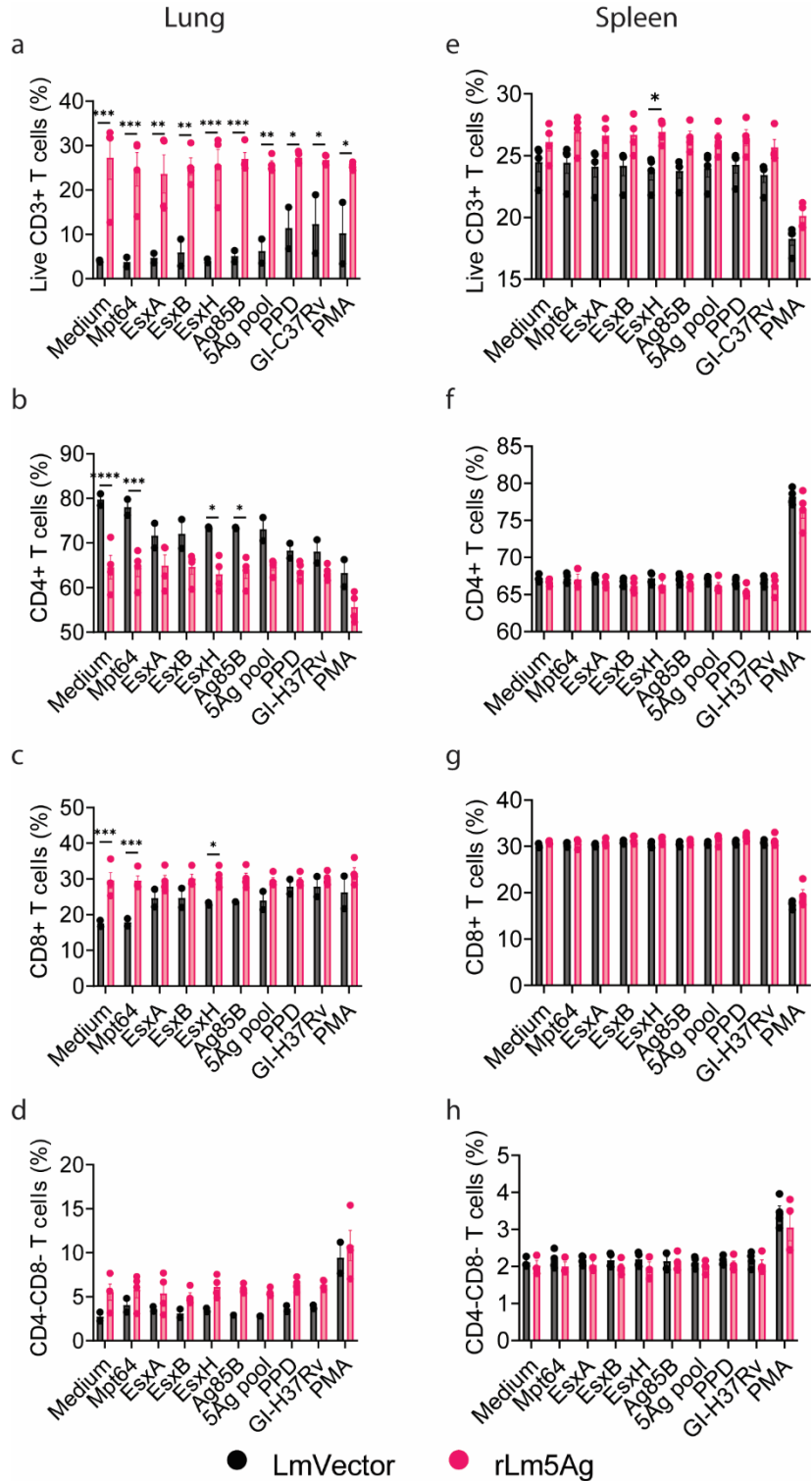
Brilliant Violet 510, BD Pharmingen) and CD8a (Clone 53-6.7, conjugated with Brilliant Violet 605, BioLegend). Afterwards, cells were fixed/permeabilized with Cytotfix/Cytoperm (BD BioSciences) and stained with antibodies to CD3 (clone 145-2C11, conjugated with Alexa Fluor 488, BD Pharmingen), IFN- $\gamma$  (Clone XMG1.2, conjugated with Brilliant Violet 711, BE Horizon), TNF- $\alpha$  (Clone MP6-XT22, conjugated with Brilliant Violet 421, BD Horizon), IL-2 (Clone JES6-5H4, conjugated with PE, BD Horizon), and IL-17A (Clone TC11-18H10.1, conjugated with Alexa Fluor 647, BD Horizon). Note that due to the internalization of CD3 in responding CD4<sup>+</sup> T cells, cells were stained for CD3 after fixing/permeabilization. A minimum of 100,000 stained cells per sample was acquired with an LSRII-HT (BD) flow cytometer and analyzed using FlowJo software (BD BioSciences). Initial gating of total events included a lymphocyte gate, followed by selection for singlet cells and live CD3<sup>+</sup> T cells and subsequently for CD4<sup>+</sup>CD8<sup>-</sup> (CD4<sup>+</sup>) and CD4<sup>-</sup>CD8<sup>+</sup> (CD8<sup>+</sup>) T cells (top panels). The gates for frequencies of antigen-specific IFN- $\gamma$ , TNF- $\alpha$ , IL-2, and IL-17A producing CD4<sup>+</sup> (middle panels) and CD8<sup>+</sup> T cells (bottom panels) were determined using the unstimulated cells. Plots shown here are spleen cells from a mouse immunized with rLm5Ag and stimulated with PPD for a total of 6 h. Pseudo colors were used to differentiate various cell populations. Frequencies of each cell population are shown within each plot. Note: In LSRII-HT (BD) flow cytometer, fixable Near-IR can be detected by the same detector as for APC-Cy7; AF488 by the same detector as for FITC; BV510 by the same detector as for Amy-Cyan; BV421 by the same detector as for Pacific blue; and AF647 by the same detector as for APC, as shown in the plots. TBm089, TB mouse experiment 89.





**Supplementary Figure 5. Frequency of antigen-specific T cells in the lungs and spleens of C57BL/6 mice immunized three times with LmVector or rLm5Ag.** C57BL/6 mice (n = 4/group, the same mice as shown in Figure 2) were immunized three times s.q. with LmVector (black) or rLm5Ag (pink) at Weeks 0, 4, and 8. At Week 9, six

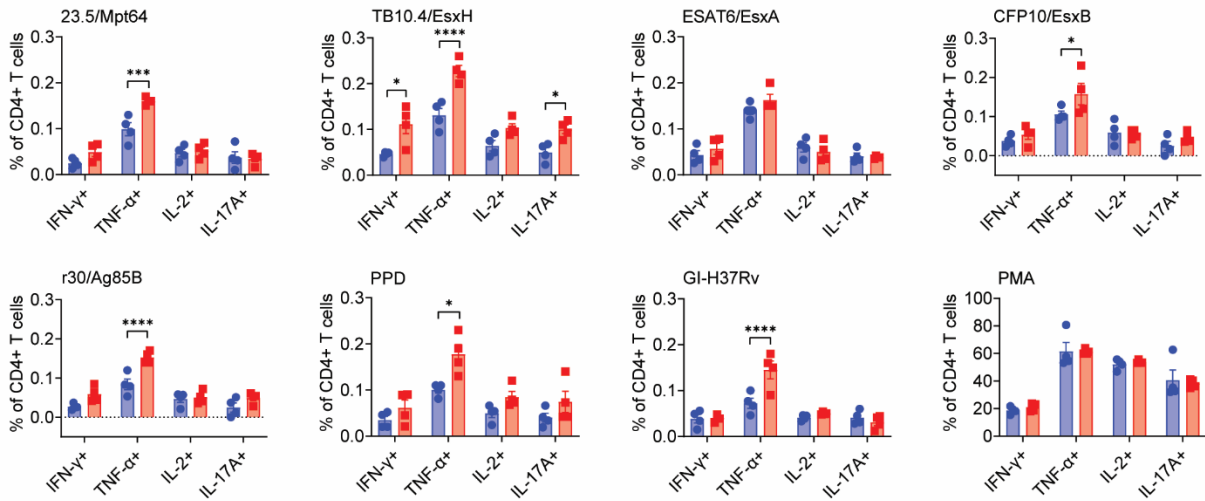
days after the last immunization, animals were euthanized; their lungs and spleens removed; and single cell suspensions prepared and stimulated in T-cell medium supplemented without (Medium) or with 2µg/ml Mpt64, EsxA, EsxB, EsxH, or Ag85B, pool of 5 antigens (5Ag pool, 2µg/ml per Ag), or PPD (2µg/ml) in the presence of anti-CD28 monoclonal antibody for a total of 22 hours, as indicated below each panel. Five hours prior to harvest, GolgiPlug (protein transport inhibitor containing Brefeldin A) diluted in T-cell medium was added to all wells; additional cells incubated with PMA and Golgiplug for 5 hours served as positive control. The cells were assayed by Live/dead staining followed by staining for surface markers of CD4, CD8 and CD3 and intracellular markers of IFN- $\gamma$ , TNF- $\alpha$ , IL-2, and IL-17A. Frequencies of live CD3+ T cells in the gated lymphocytes and frequencies of CD4+, CD8+, and CD4-CD8- T cells among live CD3+ T cells were analyzed by FlowJo 10 software. Note: frequency of CD4+CD8+ T cells among CD3+ T cells and cytokine expressing T-cells are not shown. Values are the mean  $\pm$  SEM. \*,  $P < 0.05$ ; \*\*,  $P < 0.01$ ; \*\*\*,  $P < 0.001$ ; and \*\*\*\*,  $P < 0.0001$  by two-way ANOVA with Sidak's post multiple comparisons test.



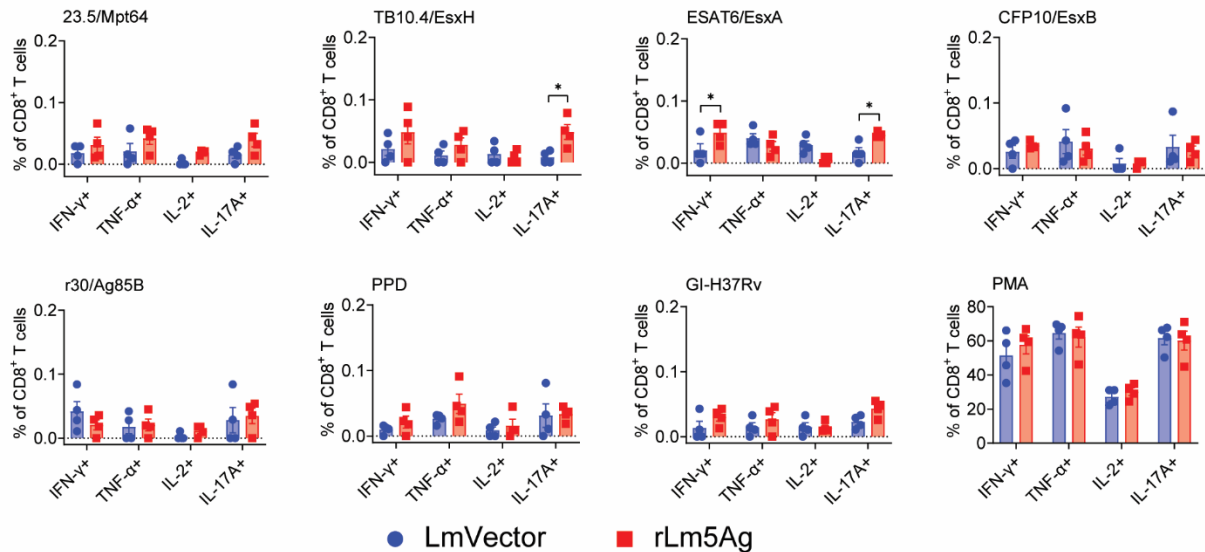
**Supplementary Figure 6. Frequency of antigen-specific T cells in the lungs and spleens of BALB/c mice immunized three times with LmVector or rLm5Ag.** BALB/c mice (n = 4/group) were immunized three times s.q. with LmVector (black) or rLm5Ag (pink) at Weeks 0, 4, and 8. At Week 9, six days after the last immunization, animals

were euthanized; their lungs and spleens removed, and single cell suspensions prepared and stimulated in T-cell medium supplemented without antigen (Medium) or with 2 $\mu$ g/ml each of Mpt64, EsxA, EsxB, EsxH, or Ag85B, pool of 5 antigens (5Ag pool, 2 $\mu$ g/ml per Ag), or PPD (2  $\mu$ g/ml) in the presence of anti-CD28 monoclonal antibody, or with gamma-irradiated Mtb H37Rv for a total of 22 hours, as indicated below each panel. Five hours prior to harvest, GolgiPlug (protein transport inhibitor containing Brefeldin A) diluted in T-cell medium was added to all wells; additional cells incubated with PMA and Golgiplug for 5 hours served as positive control. The cells were assayed by Live/dead staining followed by staining for surface markers of CD4, CD8 and CD3 and intracellular markers of IFN- $\gamma$ , TNF- $\alpha$ , IL-2, and IL-17A. Frequencies of live CD3+ T cells in the gated lymphocyte population, frequencies of CD4+, CD8+, and CD4-CD8- T cells among CD3+ T cells were analyzed by FlowJo 10 software. Note: frequency of CD4+CD8+ T cells among CD3+ T cells and cytokine producing T cells are not shown. Values are the mean  $\pm$  SEM. Each symbol represents one animal. \*,  $P < 0.05$ ; \*\*,  $P < 0.01$ ; \*\*\*,  $P < 0.001$ ; and \*\*\*\*,  $P < 0.0001$  by two-way ANOVA with Sidak's post multiple comparisons test. The experiment was repeated once in BALB/c mice.

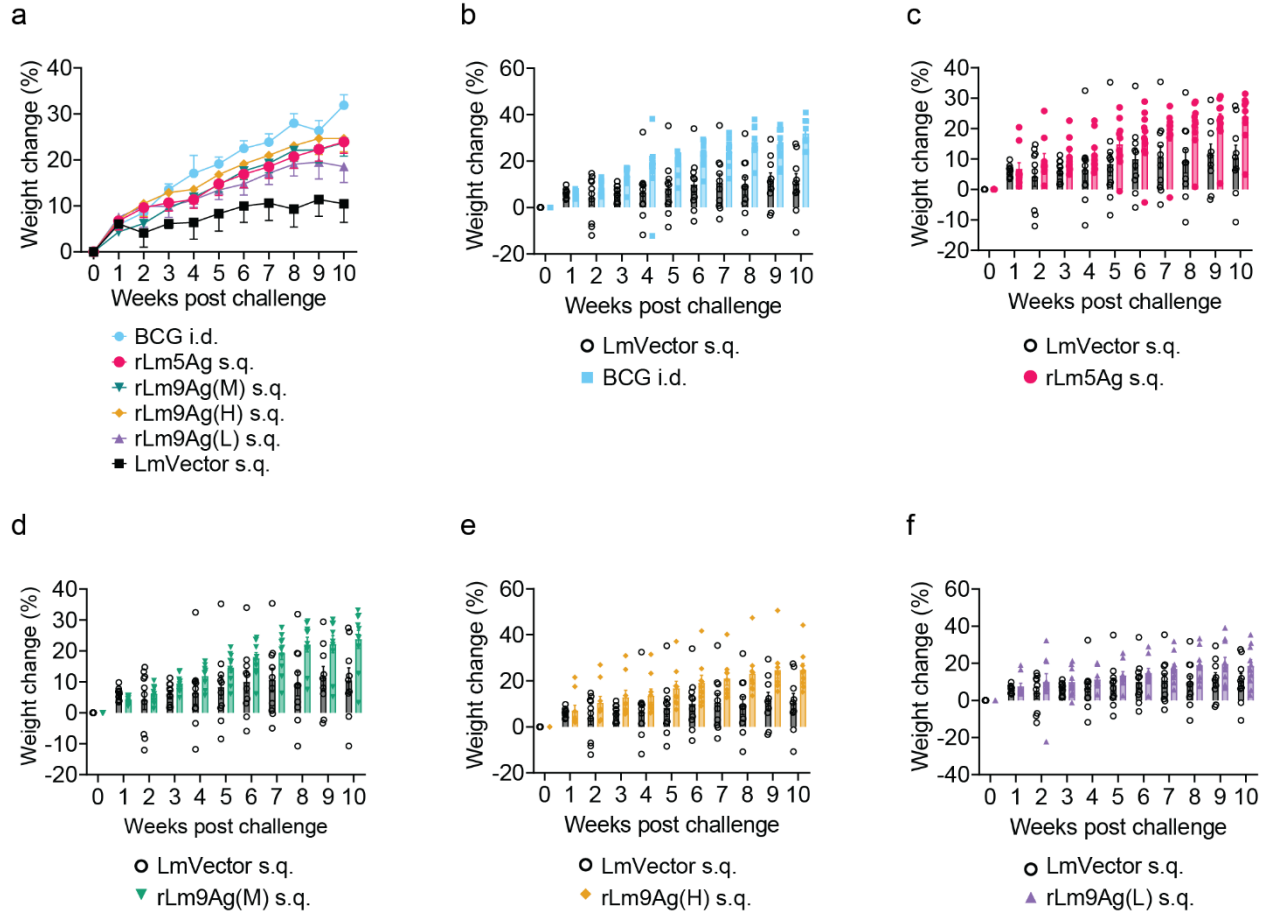
## CD4<sup>+</sup> T cells



## CD8<sup>+</sup> T cells



**Supplementary Figure 7. Frequency of cytokine-expressing CD4<sup>+</sup> and CD8<sup>+</sup> T cells in the spleens of BALB/c mice immunized three times with LmVector or rLm5Ag.** BALB/c mice (n = 4/group, the same mice as shown in Supplementary Figure 3) were immunized three times s.q. with LmVector (blue) or rLm5Ag (red) at Weeks 0, 4, and 8. One week after the last immunization, animals were euthanized, their lung and spleen cells stimulated with or without antigens for a total of 6 hours, and the cells subsequently processed as described in the legend to Supplementary Figure 3. The cells were assayed by intracellular cytokine staining for surface markers of CD4 and CD8 followed by CD3 and intracellular markers of IFN- $\gamma$ , TNF- $\alpha$ , IL-2, and IL-17A. Frequencies of splenic CD4<sup>+</sup> and CD8<sup>+</sup> T cells expressing any of the four cytokines are analyzed by FlowJo 10 software. Values are the mean  $\pm$  SEM. \*,  $P < 0.05$ ; \*\*\*,  $P < 0.001$ ; \*\*\*\*,  $P < 0.0001$  by two-way ANOVA with Sidak's post multiple comparisons test.



**Supplementary Figure 8. Data point presentation for Figure 10b. a.** The same data as shown in Fig. 10b. **b – f.** Shown are data points for panel **a** comparing vaccine group versus the LmVector group.

## Supplementary Table

**Supplementary Table 1. Amino acid sequences of Mtb antigens expressed by rLmMtb vaccine candidates**

rLm vaccine	Antigen expression cassette	Amino acid sequence
rLm30	ActA-Ag85B	VGLNRFMRAMMVVFITANCITINPDIIFAATDSEDSSLNTDEWEEEEKTEEQP SEVNTGPRYETAREVSSRDIEELEKSNKVKNTNKADLIAMLKAKAEKGGSF SRPGLPVEYLQVPSPSMGRDIKVQFQSGGNNSPAVYLLDGLRAQDDYNG WDINTPAFEWYYQSGLSIVMPVGGQSSFYSDWYSPACGKAGCQTYKWET FLTSELPQWLSANRAVKPTGSAAIGLSMAGSSAMILAAYHPQQFIYAGSLS ALLDPSQGMGPSLIGLAMGDAGGYKAADMWGPSSDPAWERNDPTQQIPK LVANNTRLWVYCGNGTPNELGGANIPAEFLENFVRSSNLKFQDAYNAAGG HNAVFNFPNGTHSWEYWGAQLNAMKGDQLQSSLGAG
rLmMtb 4Ag	ActAN-Mpt64- EsxH- GGSG- EsxA- (GSSG) <sup>2</sup> - EsxB	VGLNRFMRAMMVVFITANCITINPDIIFAATDSEDSSLNTDEWEEEEKTEEQP SEVNTGPRYETAREVSSRDIEELEKSNKVKNTNKADLIAMLKAKAEKGGSA PKTYCEELKGTDTGQACQIQMSDPAYNINISLPSYYPDQKSLENYIAQTRD KFLSAATSSTPREAPYELNITSATYQSAIPPRGTQAVVLKVYQNAGGTHPTT TYKAFDWDQAYRKPIYDYLWQADTDPLPVVFPVQVQELSKQTGQQVSIAP NAGLDPVNYQNFAVTNDGVIFFFNPGELLPEAAGPTQVLVPRSAIDSMLAR PMSQIMYNYPAMLGHAGDMAGYAGTLQSLGAEIAVEQAALQSAWQGDG ITYQAWQAQWNQAMEDLVRAYHAMSSTHEANTMAMMARDTAEAAKWG GGGSGMTEQQWNFAGIEAAASAIQGNVTSIHSLLDDEGKQSLTKLAAWGG SGSEAYQGVQKWDATATELNNALQNLARTISEAGQAMASTEGNVTGMF AGSSGGSSGMAEMKTDATLAQEAGNFERISGDLKTQIDQVESTAGSLQG QWRGAAGTAAQAAVVRFQEAANKQKQELDEISTNIRQAGVQYSRADEEQ QQALSSQMGMF
rLmMtb 5Ag	ActAN-Mpt64- EsxH- GGSG- EsxA- (GSSG) <sup>2</sup> - EsxB- (GSSG) <sup>2</sup> - Ag85B	VGLNRFMRAMMVVFITANCITINPDIIFAATDSEDSSLNTDEWEEEEKTEEQP SEVNTGPRYETAREVSSRDIEELEKSNKVKNTNKADLIAMLKAKAEKGGSM APKTYCEELKGTDTGQACQIQMSDPAYNINISLPSYYPDQKSLENYIAQTR DKFLSAATSSTPREAPYELNITSATYQSAIPPRGTQAVVLKVYQNAGGTHP TTTTYKAFDWDQAYRKPIYDYLWQADTDPLPVVFPVQVQELSKQTGQQVSI APNAGLDPVNYQNFAVTNDGVIFFFNPGELLPEAAGPTQVLVPRSAIDSML ARPMSQIMYNYPAMLGHAGDMAGYAGTLQSLGAEIAVEQAALQSAWQGD TGITYQAWQAQWNQAMEDLVRAYHAMSSTHEANTMAMMARDTAEAAKW GGGSGMTEQQWNFAGIEAAASAIQGNVTSIHSLLDDEGKQSLTKLAAWGG GSGSEAYQGVQKWDATATELNNALQNLARTISEAGQAMASTEGNVTGM FAGSSGGSSGMAEMKTDATLAQEAGNFERISGDLKTQIDQVESTAGSLQ GQWRGAAGTAAQAAVVRFQEAANKQKQELDEISTNIRQAGVQYSRADEE QQALSSQMGMFGSSGGSSGAFSRPGLPVEYLQVPSPSMGRDIKVQFQSG GNNSPAVYLLDGLRAQDDYNGWDINTPAFEWYYQSGLSIVMPVGGQSSF YSDWYSPACGKAGCQTYKWETFLTSELPQWLSANRAVKPTGSAAIGLSM AGSSAMILAAYHPQQFIYAGSLSALLDPSQGMGPSLIGLAMGDAGGYKAA DMWGPSSDPAWERNDPTQQIPKLVANNTRLWVYCGNGTPNELGGANIPA EFLENFVRSSNLKFQDAYNAAGGHNAVFNFPNGTHSWEYWGAQLNAMK GDLQSSLGAG

<p>rLmMtb 5AgII</p>	<p>ActAN- Mpt64- EsxN- (GGSG)- PPE68- (GSSG)2- EspA- (GSSG)2- TB8.4</p>	<p>VGLNRFMRAMMVVFITANCITINPDIIFAATDSEDSSLNTDEWEEEEKTEEQP SEVNTGPRYETAREVSSRDIEELEKSNKVKNTNKADLIAMLKAKAEKGGSM APKTYCEELKGTDTGQACQIQMSDPAYNINISLPSYYPDQKSLENYIAQTR DKFLSAATSSTPREAPYELNITSATYQSAIPPRGTQAVVLKVYQNAGGTHP TTTTYKAFDWDQAYRKPITYDTLWQADTDPLPVVFPVQVQVSI APNAGLDPVNYQNFVAVTNDGVIFFFNPGELLPEAAGPTQVLVPRSAIDSM AMTINYQFGDVAHGAMIRAQAASLEAEHQAVRDVLAAGDFWGGAGSVA CQEFITQLGRNFQVIYEQANAHGQKVQAAGNNMAQTD SAVGSSWAGGSG MLWHAMPPELNTARLMAGAGPAPMLAAAAGWQTL SAALDAQAVELTARL NSLGEAWTGGGSDKALAAATPMVVWLQTASTQAKTRAMQATAQAAAAYTQ AMATTPSLPEIAANHITQAVLTATNFFGINTIPIALTEMDYFIRMWNQAALAM EVYQAETAVNTLFEKLEPMASILDPGASQSTTNP IFGMPSPGSSSTPVGQLP PAATQTLGQLGEMSGPMQQLTQPLQQVTSLFSQVGGTGGGNPADEEAA QMGLLGTSPLSNHPLAGGSGPSAGAGLLRAESLPGAGGSLTRTPLMSQLI EKPVAPSVMPAAAAGSSATGGAAPVGAGAMGQGAQSGGSTRPGLVAPA PLAQEREEDDEDDWDEEDDWGSSGGSSGAMSRAFIIDPTISAIDGLYDLL GIGIPNQGGILYSSLEYFEKALEELAAAFPGDGWLGSAADKYAGKNRNHVN FFQELADLDRQLISLIHDQANAVQTTRDILEGAKKGLEGEVWEFITNALNGL KELWDKLTGWVTGLFSRGSWNLESFFAGVPGLTGATSGLSQVTGLFGAA GLSASSGLAHADSLASSASLPALAGIGGGSGFGGLPSLAQVHAASRQAL RPRADGPVGA AA EQVGGQSQLVSAQGSQGMGGPVGMGGMHPSSGASK GTTTKKYSEGAAAGTEDAERAPVEADAGGGQKVLVRNVVSSGGSSGAM DPVDAVINTTCNYGQVVAALNATDPGAAAQFNASPV AQSYLRNFLAAPP QRAAMAAQLQAVPGAAQYIGLVESVAGSCNNY</p>
<p>rLmMtb 9Ag</p>	<p>ActAN- Mpt64- EsxH- GGSG- EsxA- (GSSG)2- EsxB- (GSSG2)- Ag85B</p>	<p>The same as for rLm5Ag</p>
<p>rLmMtb 9Ag</p>	<p>ActAN- Mpt64- EsxN- (GGSG)- PPE68- (GSSG)2- EspA- (GSSG)2- TB8.4</p>	<p>The same as for rLm5AgII</p>



**Supplementary Table 2. Statistical analyses comparing weight change post challenge in guinea pig groups by 2-Way ANOVA with Tukey's multiple comparisons test (Prism)**

Tukey's multiple comparisons test	95.00% CI of diff.	Below threshold?	Summary	Adjusted P Value
<b>0 Week post challenge</b>				
BCG i.d. vs. LmVector s.q.	-9.912 to 9.912	No	ns	>0.9999
BCG i.d. vs. rLm9Ag(L) s.q.	-9.912 to 9.912	No	ns	>0.9999
BCG i.d. vs. rLm9Ag(M) s.q.	-9.912 to 9.912	No	ns	>0.9999
BCG i.d. vs. rLm9Ag(H) s.q.	-10.18 to 10.18	No	ns	>0.9999
BCG i.d. vs. rLm5Ag s.q.	-9.912 to 9.912	No	ns	>0.9999
LmVector s.q. vs. rLm9Ag(L) s.q.	-9.912 to 9.912	No	ns	>0.9999
LmVector s.q. vs. rLm9Ag(M) s.q.	-9.912 to 9.912	No	ns	>0.9999
LmVector s.q. vs. rLm9Ag(H) s.q.	-10.18 to 10.18	No	ns	>0.9999
LmVector s.q. vs. rLm5Ag s.q.	-9.912 to 9.912	No	ns	>0.9999
rLm9Ag(L) s.q. vs. rLm9Ag(M) s.q.	-9.912 to 9.912	No	ns	>0.9999
rLm9Ag(L) s.q. vs. rLm9Ag(H) s.q.	-10.18 to 10.18	No	ns	>0.9999
rLm9Ag(L) s.q. vs. rLm5Ag s.q.	-9.912 to 9.912	No	ns	>0.9999
rLm9Ag(M) s.q. vs. rLm9Ag(H) s.q.	-10.18 to 10.18	No	ns	>0.9999
rLm9Ag(M) s.q. vs. rLm5Ag s.q.	-9.912 to 9.912	No	ns	>0.9999
rLm9Ag(H) s.q. vs. rLm5Ag s.q.	-10.18 to 10.18	No	ns	>0.9999
<b>1 Week post challenge</b>				
BCG i.d. vs. LmVector s.q.	-10.09 to 9.734	No	ns	>0.9999
BCG i.d. vs. rLm9Ag(L) s.q.	-11.51 to 8.309	No	ns	0.9974
BCG i.d. vs. rLm9Ag(M) s.q.	-8.348 to 11.48	No	ns	0.9976
BCG i.d. vs. rLm9Ag(H) s.q.	-11.32 to 9.047	No	ns	0.9996
BCG i.d. vs. rLm5Ag s.q.	-10.74 to 9.080	No	ns	0.9999
LmVector s.q. vs. rLm9Ag(L) s.q.	-11.34 to 8.487	No	ns	0.9985
LmVector s.q. vs. rLm9Ag(M) s.q.	-8.170 to 11.65	No	ns	0.9961
LmVector s.q. vs. rLm9Ag(H) s.q.	-11.14 to 9.225	No	ns	0.9998
LmVector s.q. vs. rLm5Ag s.q.	-10.57 to 9.258	No	ns	>0.9999
rLm9Ag(L) s.q. vs. rLm9Ag(M) s.q.	-6.745 to 13.08	No	ns	0.943
rLm9Ag(L) s.q. vs. rLm9Ag(H) s.q.	-9.717 to 10.65	No	ns	>0.9999
rLm9Ag(L) s.q. vs. rLm5Ag s.q.	-9.141 to 10.68	No	ns	>0.9999
rLm9Ag(M) s.q. vs. rLm9Ag(H) s.q.	-12.88 to 7.483	No	ns	0.9742
rLm9Ag(M) s.q. vs. rLm5Ag s.q.	-12.31 to 7.516	No	ns	0.9829
rLm9Ag(H) s.q. vs. rLm5Ag s.q.	-9.879 to 10.49	No	ns	>0.9999
<b>2 Weeks post challenge</b>				
BCG i.d. vs. LmVector s.q.	-5.470 to 14.35	No	ns	0.7952
BCG i.d. vs. rLm9Ag(L) s.q.	-11.29 to 8.537	No	ns	0.9987
BCG i.d. vs. rLm9Ag(M) s.q.	-7.534 to 12.29	No	ns	0.9835

BCG i.d. vs. rLm9Ag(H) s.q.	-12.14 to 8.225	No	ns	0.994
BCG i.d. vs. rLm5Ag s.q.	-10.99 to 8.830	No	ns	0.9996
LmVector s.q. vs. rLm9Ag(L) s.q.	-15.73 to 4.095	No	ns	0.5469
LmVector s.q. vs. rLm9Ag(M) s.q.	-11.98 to 7.848	No	ns	0.9913
LmVector s.q. vs. rLm9Ag(H) s.q.	-16.58 to 3.783	No	ns	0.4685
LmVector s.q. vs. rLm5Ag s.q.	-15.44 to 4.388	No	ns	0.6032
rLm9Ag(L) s.q. vs. rLm9Ag(M) s.q.	-6.159 to 13.66	No	ns	0.8883
rLm9Ag(L) s.q. vs. rLm9Ag(H) s.q.	-10.77 to 9.600	No	ns	>0.9999
rLm9Ag(L) s.q. vs. rLm5Ag s.q.	-9.619 to 10.20	No	ns	>0.9999
rLm9Ag(M) s.q. vs. rLm9Ag(H) s.q.	-14.52 to 5.847	No	ns	0.8283
rLm9Ag(M) s.q. vs. rLm5Ag s.q.	-13.37 to 6.452	No	ns	0.9185
rLm9Ag(H) s.q. vs. rLm5Ag s.q.	-9.307 to 11.06	No	ns	0.9999
<b>3 Weeks post challenge</b>				
BCG i.d. vs. LmVector s.q.	-2.493 to 17.33	No	ns	0.2681
BCG i.d. vs. rLm9Ag(L) s.q.	-6.204 to 13.62	No	ns	0.8933
BCG i.d. vs. rLm9Ag(M) s.q.	-5.802 to 14.02	No	ns	0.8437
BCG i.d. vs. rLm9Ag(H) s.q.	-9.508 to 10.86	No	ns	>0.9999
BCG i.d. vs. rLm5Ag s.q.	-7.003 to 12.82	No	ns	0.9601
LmVector s.q. vs. rLm9Ag(L) s.q.	-13.62 to 6.201	No	ns	0.8929
LmVector s.q. vs. rLm9Ag(M) s.q.	-13.22 to 6.603	No	ns	0.9318
LmVector s.q. vs. rLm9Ag(H) s.q.	-16.93 to 3.439	No	ns	0.4071
LmVector s.q. vs. rLm5Ag s.q.	-14.42 to 5.402	No	ns	0.7845
rLm9Ag(L) s.q. vs. rLm9Ag(M) s.q.	-9.510 to 10.31	No	ns	>0.9999
rLm9Ag(L) s.q. vs. rLm9Ag(H) s.q.	-13.22 to 7.150	No	ns	0.9575
rLm9Ag(L) s.q. vs. rLm5Ag s.q.	-10.71 to 9.113	No	ns	>0.9999
rLm9Ag(M) s.q. vs. rLm9Ag(H) s.q.	-13.62 to 6.748	No	ns	0.9289
rLm9Ag(M) s.q. vs. rLm5Ag s.q.	-11.11 to 8.711	No	ns	0.9993
rLm9Ag(H) s.q. vs. rLm5Ag s.q.	-7.949 to 12.42	No	ns	0.989
<b>4 Weeks post challenge</b>				
BCG i.d. vs. LmVector s.q.	0.7654 to 20.59	Yes	*	0.0263
BCG i.d. vs. rLm9Ag(L) s.q.	-4.151 to 15.67	No	ns	0.5577
BCG i.d. vs. rLm9Ag(M) s.q.	-4.729 to 15.09	No	ns	0.6675
BCG i.d. vs. rLm9Ag(H) s.q.	-6.693 to 13.67	No	ns	0.9242
BCG i.d. vs. rLm5Ag s.q.	-4.190 to 15.63	No	ns	0.5652
LmVector s.q. vs. rLm9Ag(L) s.q.	-14.83 to 4.996	No	ns	0.716
LmVector s.q. vs. rLm9Ag(M) s.q.	-15.41 to 4.418	No	ns	0.6089
LmVector s.q. vs. rLm9Ag(H) s.q.	-17.37 to 2.997	No	ns	0.3334
LmVector s.q. vs. rLm5Ag s.q.	-14.87 to 4.957	No	ns	0.7091
rLm9Ag(L) s.q. vs. rLm9Ag(M) s.q.	-10.49 to 9.334	No	ns	>0.9999
rLm9Ag(L) s.q. vs. rLm9Ag(H) s.q.	-12.45 to 7.913	No	ns	0.9881
rLm9Ag(L) s.q. vs. rLm5Ag s.q.	-9.951 to 9.873	No	ns	>0.9999
rLm9Ag(M) s.q. vs. rLm9Ag(H) s.q.	-11.88 to 8.491	No	ns	0.997

rLm9Ag(M) s.q. vs. rLm5Ag s.q.	-9.373 to 10.45	No	ns	>0.9999
rLm9Ag(H) s.q. vs. rLm5Ag s.q.	-7.952 to 12.41	No	ns	0.989
5 Weeks post challenge				
BCG i.d. vs. LmVector s.q.	0.9304 to 20.75	Yes	*	0.0227
BCG i.d. vs. rLm9Ag(L) s.q.	-4.207 to 15.62	No	ns	0.5684
BCG i.d. vs. rLm9Ag(M) s.q.	-5.325 to 14.50	No	ns	0.7721
BCG i.d. vs. rLm9Ag(H) s.q.	-7.860 to 12.51	No	ns	0.9868
BCG i.d. vs. rLm5Ag s.q.	-5.551 to 14.27	No	ns	0.8076
LmVector s.q. vs. rLm9Ag(L) s.q.	-15.05 to 4.775	No	ns	0.676
LmVector s.q. vs. rLm9Ag(M) s.q.	-16.17 to 3.657	No	ns	0.4636
LmVector s.q. vs. rLm9Ag(H) s.q.	-18.70 to 1.665	No	ns	0.1607
LmVector s.q. vs. rLm5Ag s.q.	-16.39 to 3.431	No	ns	0.422
rLm9Ag(L) s.q. vs. rLm9Ag(M) s.q.	-11.03 to 8.794	No	ns	0.9995
rLm9Ag(L) s.q. vs. rLm9Ag(H) s.q.	-13.56 to 6.802	No	ns	0.9333
rLm9Ag(L) s.q. vs. rLm5Ag s.q.	-11.26 to 8.568	No	ns	0.9989
rLm9Ag(M) s.q. vs. rLm9Ag(H) s.q.	-12.45 to 7.920	No	ns	0.9883
rLm9Ag(M) s.q. vs. rLm5Ag s.q.	-10.14 to 9.686	No	ns	>0.9999
rLm9Ag(H) s.q. vs. rLm5Ag s.q.	-8.146 to 12.22	No	ns	0.9928
6 Weeks post challenge				
BCG i.d. vs. LmVector s.q.	2.643 to 22.47	Yes	**	0.0043
BCG i.d. vs. rLm9Ag(L) s.q.	-2.163 to 17.66	No	ns	0.2231
BCG i.d. vs. rLm9Ag(M) s.q.	-4.993 to 14.83	No	ns	0.7155
BCG i.d. vs. rLm9Ag(H) s.q.	-6.808 to 13.56	No	ns	0.9338
BCG i.d. vs. rLm5Ag s.q.	-4.286 to 15.54	No	ns	0.5836
LmVector s.q. vs. rLm9Ag(L) s.q.	-14.72 to 5.106	No	ns	0.7353
LmVector s.q. vs. rLm9Ag(M) s.q.	-17.55 to 2.276	No	ns	0.2379
LmVector s.q. vs. rLm9Ag(H) s.q.	-19.36 to 1.003	No	ns	0.1044
LmVector s.q. vs. rLm5Ag s.q.	-16.84 to 2.983	No	ns	0.3442
rLm9Ag(L) s.q. vs. rLm9Ag(M) s.q.	-12.74 to 7.082	No	ns	0.9645
rLm9Ag(L) s.q. vs. rLm9Ag(H) s.q.	-14.56 to 5.809	No	ns	0.823
rLm9Ag(L) s.q. vs. rLm5Ag s.q.	-12.03 to 7.789	No	ns	0.9901
rLm9Ag(M) s.q. vs. rLm9Ag(H) s.q.	-11.73 to 8.639	No	ns	0.9981
rLm9Ag(M) s.q. vs. rLm5Ag s.q.	-9.205 to 10.62	No	ns	>0.9999
rLm9Ag(H) s.q. vs. rLm5Ag s.q.	-7.932 to 12.43	No	ns	0.9886
7 Weeks post challenge				
BCG i.d. vs. LmVector s.q.	3.371 to 23.19	Yes	**	0.002
BCG i.d. vs. rLm9Ag(L) s.q.	-3.144 to 16.68	No	ns	0.3713
BCG i.d. vs. rLm9Ag(M) s.q.	-5.288 to 14.54	No	ns	0.766
BCG i.d. vs. rLm9Ag(H) s.q.	-7.307 to 13.06	No	ns	0.9661
BCG i.d. vs. rLm5Ag s.q.	-4.561 to 15.26	No	ns	0.6361
LmVector s.q. vs. rLm9Ag(L) s.q.	-16.43 to 3.397	No	ns	0.4159

LmVector s.q. vs. rLm9Ag(M) s.q.	-18.57 to 1.253	No	ns	0.1263
LmVector s.q. vs. rLm9Ag(H) s.q.	-20.59 to -0.2233	Yes	*	0.0419
LmVector s.q. vs. rLm5Ag s.q.	-17.84 to 1.980	No	ns	0.2005
rLm9Ag(L) s.q. vs. rLm9Ag(M) s.q.	-12.06 to 7.768	No	ns	0.9897
rLm9Ag(L) s.q. vs. rLm9Ag(H) s.q.	-14.07 to 6.292	No	ns	0.8843
rLm9Ag(L) s.q. vs. rLm5Ag s.q.	-11.33 to 8.495	No	ns	0.9985
rLm9Ag(M) s.q. vs. rLm9Ag(H) s.q.	-11.93 to 8.436	No	ns	0.9965
rLm9Ag(M) s.q. vs. rLm5Ag s.q.	-9.185 to 10.64	No	ns	>0.9999
rLm9Ag(H) s.q. vs. rLm5Ag s.q.	-7.709 to 12.66	No	ns	0.9825
<b>8 Weeks post challenge</b>				
BCG i.d. vs. LmVector s.q.	8.769 to 28.59	Yes	****	<0.0001
BCG i.d. vs. rLm9Ag(L) s.q.	-1.004 to 18.82	No	ns	0.1064
BCG i.d. vs. rLm9Ag(M) s.q.	-4.000 to 15.82	No	ns	0.5287
BCG i.d. vs. rLm9Ag(H) s.q.	-5.308 to 15.06	No	ns	0.7457
BCG i.d. vs. rLm5Ag s.q.	-2.637 to 17.19	No	ns	0.2893
LmVector s.q. vs. rLm9Ag(L) s.q.	-19.68 to 0.1386	No	ns	0.0558
LmVector s.q. vs. rLm9Ag(M) s.q.	-22.68 to -2.857	Yes	**	0.0034
LmVector s.q. vs. rLm9Ag(H) s.q.	-23.99 to -3.622	Yes	**	0.0016
LmVector s.q. vs. rLm5Ag s.q.	-21.32 to -1.494	Yes	*	0.0135
rLm9Ag(L) s.q. vs. rLm9Ag(M) s.q.	-12.91 to 6.916	No	ns	0.9548
rLm9Ag(L) s.q. vs. rLm9Ag(H) s.q.	-14.22 to 6.151	No	ns	0.8678
rLm9Ag(L) s.q. vs. rLm5Ag s.q.	-11.54 to 8.279	No	ns	0.9971
rLm9Ag(M) s.q. vs. rLm9Ag(H) s.q.	-11.22 to 9.147	No	ns	0.9997
rLm9Ag(M) s.q. vs. rLm5Ag s.q.	-8.549 to 11.27	No	ns	0.9988
rLm9Ag(H) s.q. vs. rLm5Ag s.q.	-7.784 to 12.58	No	ns	0.9848
<b>9 Weeks post challenge</b>				
BCG i.d. vs. LmVector s.q.	4.797 to 25.16	Yes	***	0.0004
BCG i.d. vs. rLm9Ag(L) s.q.	-3.093 to 16.73	No	ns	0.3626
BCG i.d. vs. rLm9Ag(M) s.q.	-5.738 to 14.09	No	ns	0.8349
BCG i.d. vs. rLm9Ag(H) s.q.	-8.488 to 11.88	No	ns	0.997
BCG i.d. vs. rLm5Ag s.q.	-5.863 to 13.96	No	ns	0.8519
LmVector s.q. vs. rLm9Ag(L) s.q.	-18.34 to 2.022	No	ns	0.1991
LmVector s.q. vs. rLm9Ag(M) s.q.	-20.99 to -0.6227	Yes	*	0.0302
LmVector s.q. vs. rLm9Ag(H) s.q.	-23.73 to -2.837	Yes	**	0.0041
LmVector s.q. vs. rLm5Ag s.q.	-21.11 to -0.7477	Yes	*	0.0271
rLm9Ag(L) s.q. vs. rLm9Ag(M) s.q.	-12.56 to 7.267	No	ns	0.9735
rLm9Ag(L) s.q. vs. rLm9Ag(H) s.q.	-15.31 to 5.060	No	ns	0.7033
rLm9Ag(L) s.q. vs. rLm5Ag s.q.	-12.68 to 7.142	No	ns	0.9676
rLm9Ag(M) s.q. vs. rLm9Ag(H) s.q.	-12.66 to 7.705	No	ns	0.9824
rLm9Ag(M) s.q. vs. rLm5Ag s.q.	-10.04 to 9.787	No	ns	>0.9999
rLm9Ag(H) s.q. vs. rLm5Ag s.q.	-7.830 to 12.54	No	ns	0.986

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10 Weeks post challenge

BCG i.d. vs. LmVector s.q.	10.68 to 32.21	Yes	****	<0.0001
BCG i.d. vs. rLm9Ag(L) s.q.	2.850 to 23.88	Yes	**	0.0041
BCG i.d. vs. rLm9Ag(M) s.q.	-2.388 to 18.64	No	ns	0.2347
BCG i.d. vs. rLm9Ag(H) s.q.	-3.531 to 18.01	No	ns	0.3898
BCG i.d. vs. rLm5Ag s.q.	-2.536 to 18.49	No	ns	0.2536
LmVector s.q. vs. rLm9Ag(L) s.q.	-18.26 to 2.102	No	ns	0.2085
LmVector s.q. vs. rLm9Ag(M) s.q.	-23.50 to -3.136	Yes	**	0.0028
LmVector s.q. vs. rLm9Ag(H) s.q.	-24.65 to -3.759	Yes	**	0.0016
LmVector s.q. vs. rLm5Ag s.q.	-23.65 to -3.284	Yes	**	0.0024
rLm9Ag(L) s.q. vs. rLm9Ag(M) s.q.	-15.15 to 4.674	No	ns	0.6573
rLm9Ag(L) s.q. vs. rLm9Ag(H) s.q.	-16.31 to 4.058	No	ns	0.5191
rLm9Ag(L) s.q. vs. rLm5Ag s.q.	-15.30 to 4.526	No	ns	0.6294
rLm9Ag(M) s.q. vs. rLm9Ag(H) s.q.	-11.07 to 9.296	No	ns	0.9999
rLm9Ag(M) s.q. vs. rLm5Ag s.q.	-10.06 to 9.764	No	ns	>0.9999
rLm9Ag(H) s.q. vs. rLm5Ag s.q.	-9.444 to 10.92	No	ns	>0.9999

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