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Single-dose, virus-vectored vaccine protection against *Yersinia pestis* challenge: CD4⁺ cells are required at the time of challenge for optimal protection

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Abstract

We have developed an experimental recombinant vesicular stomatitis virus (VSV) vectored plague vaccine expressing a secreted form of *Yersinia pestis* LcrV protein from the first position of the VSV genome. This vector, given intramuscularly in a single dose, induced high-level antibody titers to LcrV and gave 90–100% protection against pneumonic plague challenge in mice. This single-dose protection was significantly better than that generated by VSV expressing the non-secreted LcrV protein. Increased protection correlated with increased anti-LcrV antibody and a bias toward IgG2a and away from IgG1 isotypes. We also found that depletion of CD4⁺ cells, but not CD8⁺ cells, at the time of challenge resulted in reduced vaccine protection, indicating a role for cellular immunity in protection.

Keywords

Yersinia pestis; Vesicular stomatitis virus; CD4⁺ cells

1. Introduction

Yersinia pestis, a gram-negative bacillus that primarily causes disease of small rodents, can occasionally also infect humans via infected rodent or flea bites, thereby causing bubonic plague [1]. Untreated bubonic plague can progress to a highly contagious pneumonic form of the disease that is associated with high mortality. Humans can also contract pneumonic plague by inhalation of aerosolized infectious droplets following direct contact with an infected individual, thus allowing for the rapid spread of the disease [1,2]. With antibiotic treatments being only marginally effective [1], the rapidly progressing and often-fatal nature of pneumonic plague poses a danger to public health worldwide. In addition, the possible intentional use of aerosolized *Y. pestis* or its antibiotic-resistant forms [3,4] as bioterrorism agents looms as a threat [5].

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Vaccination against plague is one approach being pursued to deal with impending infection. However, a safe and effective pneumonic plague vaccine for human is not yet available in the United States [6,7]. Current vaccine development strategies focus on a few *Y. pestis* antigens [7], two major candidates being the outer capsule protein (F1) and the low calcium response protein V (LcrV), which induce good protective immune response against challenge with the pathogen [8,9]. While vaccines based on F1 alone fail to protect against F1⁻ virulent strains [10], LcrV based sub-unit vaccines confer protection against both F1⁺ and F1⁻ *Y. pestis* strains [10–18]. LcrV is a 37 kDa protein essential for virulence and critical for production and translocation of the *Yersinia* outer proteins (Yops) into the eukaryotic cell cytosol [19,20]. Although the mechanism of protection elicited by LcrV-based vaccines is poorly understood, a number of studies have shown that antibodies to LcrV, generated by either active or passive immunization, can protect from the disease in mouse models [21–26]. Recently, induction of LcrV-specific broad T-cell immunity has also been reported [27,28], but its role in protection remains to be elucidated.

We had previously reported the development of an experimental pneumonic plague vaccine based on an attenuated recombinant vesicular stomatitis virus (rVSV) vector expressing *Yersinia pestis* LcrV protein [24]. VSV is a negative-strand RNA virus encoding five structural proteins - nucleocapsid (N), phosphoprotein (P), matrix (M), glycoprotein (G) and RNA dependent RNA polymerase (L). The virus is a natural pathogen of livestock, and human infection is rare and typically asymptomatic [29]. VSV recombinants, that are attenuated for pathogenesis in mice compared to naturally occurring VSV strains [30], can be generated from plasmid DNA [31,32] and also cause no disease symptoms in non-human primates when given by intranasal (IN), intramuscular (IM) or oral routes [33–35]. Furthermore, VSV can accommodate large insertions of foreign genes whose expression can be controlled based on the site of gene insertion in the VSV genome [36,37]. Such recombinants induce potent humoral and cellular immune responses in a variety of animal models [29,38,39]. Numerous studies have shown that recombinant VSV vectors expressing appropriate foreign antigens are highly effective experimental vaccines [24,30,35,40–50].

In our earlier study, we showed that rVSV vector expressing the *Yersinia pestis* *lcrV* gene conferred high-level and long-term protection against intranasal plague challenge in mice, after a single booster immunization [24]. A previous study using a multiple-dose DNA vaccine found that the secreted form of LcrV induced improved antibody responses and better protection of mice from challenge [26]. In the present study, we have incorporated a gene encoding a secreted form of LcrV in the VSV vector (VSV-ssLcrV) and tested its efficacy as a single-dose vaccine against lethal plague challenge. The vector elicits robust antibody responses with increasing immunization dose in mice and confers complete protection against pulmonary *Y. pestis* challenge. The presence of vaccine induced CD4⁺ cells at the time of challenge is important for generating complete protection.

2. Materials and methods

2.1. Construction of plasmids and virus recovery

The plasmid pVSV-ssLcrV1 used to derive a VSV recombinant expressing a secreted form of low calcium response protein V (LcrV) of *Yersinia pestis* was generated as follows. The ssLcrV coding region, containing a signal sequence (from the human plasminogen activator protein) upstream of the LcrV open reading frame (ORF), was amplified by PCR using the pBS-LcrV plasmid [24] as a template. The signal sequence was added upstream of the LcrV ORF in the forward primer. The PCR was performed using Vent DNA polymerase (New England Biolabs), forward primer 5'-GATCGATC GTCGACAACATGGATGCAATGAA
GAGAGGGCTCTGCTGTGTGCTGCTGCTG TGTGGAGCAGTCTTCGT
TTCGATTAGAGCCTACGAACAAAACCC -3' [sequence encoding signal (underlined)]

followed by 2nd codon of LcrV] and reverse primer 5'-CGA TCCCCCGGGCTAGCTCATTACCAGACGTGTCATCTA GCAG -3'. The forward and the reverse primers respectively introduced *Sal* I and *Nhe* I restriction enzyme sites (in bold) upstream and downstream of the ssLcrV coding sequence. The PCR product was digested with *Sal* I and *Nhe* I, purified and ligated to *Xho* I-*Nhe* I digested pVSV1XN [51] to generate pVSV-ssLcrV1. Recombinant VSV expressing secreted LcrV from the first position in the VSV genome was recovered from these plasmids as described previously [31]. Briefly, Baby Hamster Kidney (BHK-21) cells were grown to 70% confluency and infected with vTF7-3 [52], a vaccinia virus recombinant expressing T7 RNA polymerase, at a multiplicity of infection (MOI) of 10 for 1 hr followed by transfection with pVSV-ssLcrV1 along with the support plasmids pBS-N, pBS-P and pBS-L. At 48 hrs post transfection, cell supernatants were passaged onto fresh BHK-21 cells. Medium containing the virus was collected after 48 hrs. Virus stocks were prepared from individual plaques grown in BHK-21 cells and were stored at -70°C

2.2. Indirect Immunofluorescence

BHK-21 cells seeded on glass cover slips were infected with VSV-ssLcrV1, VSV-LcrV1 or recombinant wild-type (rwt)-VSV. At 3 hrs post infection, cells were washed twice with phosphate buffered saline (PBS) and fixed with 3% Para formaldehyde at room temperature. Following this, cells were washed with PBS containing 10mM glycine (PBS-glycine) and permeabilized with 1% Triton-x-100 in PBS-glycine. Cells were then incubated with an equal mixture of 1:200 diluted anti-LcrV monoclonal antibodies (MAbs) - Va13 and Va48 (Santa Cruz Biotechnology, CA), followed by incubation with goat anti-mouse Alexa Flour 594 IgG (Molecular probes, Eugene, OR) diluted 1:500. The cells were then washed twice with PBS-glycine, mounted on slides and imaged using a Nikon Eclipse 80i fluorescence microscope equipped with a Nikon Plan APOchromat 60X oil objective and a Photometrics CoolSnap camera.

2.3. Metabolic labeling and SDS-PAGE of cells infected with recombinant viruses

BHK-21 cells were infected with recombinant viruses at an MOI of 100. After 4 hrs, cells were washed with methionine-free Dulbecco's modified Eagle's medium (DMEM) followed by labeling with 100 µCi of [³⁵S]-methionine for 2 hrs. Cell supernatants were then collected and infected cells were washed with PBS and lysed with detergent lysis buffer (1% nonidet P-40, 0.4% deoxycholate, 50mM Tris-HCl, pH8.0, 62.5 mM EDTA) on ice for 5 min to prepare lysates. Both lysates and supernatants were clarified at 16,000 g for 2 min at 4°C and then analyzed by 10% SDS-PAGE. Protein bands were visualized by autoradiography.

2.4. Vaccination of mice

Six to eight week-old Balb/C female mice were obtained from either Charles River Laboratories or Taconic Laboratories and then rested for a minimum of 1 week before inoculation. Animals were housed under BSL-2 conditions in microisolator cages. Viruses were diluted in serum free DMEM. Mice were vaccinated intramuscularly (IM) into the left hind leg muscle with doses of virus as indicated in the text in a total volume of 50µL. Immunizations performed at Yale University, and immunizations and challenges carried out at the Public Health Research Institute (PHRI), Newark, NJ, were approved by the respective Institutional Animal Care and Use Committees.

2.5. ELISA for total serum antibody to LcrV

Serum samples were heat inactivated at 56°C for 30 min. An enzyme linked immunosorbent assay (ELISA) to determine the total serum antibody specific to LcrV was performed as described by Palin *et al.* [24], with minor modifications. Briefly, 100 ng of purified GST-LcrV

protein [from Northeast Biodefense Center (NBC) Protein Expression Core, Wadsworth Center, NY] in 100 μ L capture buffer (20 mM Tris-HCl, pH 8.5 and 0.1 M NaCl) was added to each well of a 96-well plate (Corning) for 16 hrs at 4°C. Plates were blocked with 200 μ L 10% bovine serum in PBS for 30 min. 2-fold serially diluted serum samples (in duplicate) were added to the plates and incubated for 1.5 hrs at room temperature. Goat-anti-mouse secondary antibody conjugated to horseradish peroxidase (HRP) was added at a dilution of 1:10,000 (100 μ L/well) and incubated for further 1.5 hrs. Immunopure ABTS tablet (Pierce), dissolved in 10 mL substrate buffer (0.1 M acetate, 0.05% Tween 20, pH 4.2) to which 5 μ L 30% H₂O₂ was added, was used as the substrate solution (100 μ L/well) and plates were read at 415 nm on a Biorad microplate reader at 20 min following addition of substrate. Titers were calculated as the reciprocal of the maximum serum dilution giving an OD₄₁₅ of 0.2, after subtracting the non-specific binding in serum of VSV-eGFP1, a recombinant VSV expressing enhanced Green Fluorescence Protein (eGFP) from the first position [51], infected animals.

2.6. Isotype determination of LcrV specific antibodies

ELISA assays to determine the levels of IgG1 and IgG2a bound specifically to LcrV were performed as described above (section 2.5), except using HRP-conjugated goat anti-mouse IgG1 (used at 1:5,000 dilution) or IgG2a (used at 1:10,000 dilution). To calculate the amounts of IgG1 or IgG2a bound to LcrV, we also generated standard curves using a range of concentrations of mouse IgG1 or IgG2a (Sigma) bound directly to the ELISA plate and detected with HRP-conjugated goat anti-mouse IgG1 or or IgG2a (Sigma).

2.7. Depletion of CD4⁺ and/or CD8⁺ cells

To determine the extent of CD4⁺ and/or CD8⁺ cell depletion by MAbs, female Balb/C mice, obtained from Charles River laboratories were rested for 1 week before administration of 500 μ g of rat anti-mouse MAbs specific for mouse CD4 (clone GK1.5) and/or mouse CD8 (clone 2.43) by the intraperitoneal (IP) route. Control animals were either injected with 500 μ g of the isotype control Rat IgG2b (clone LTF-2) or left untreated (naïve). All MAbs were purchased from Bio Express Cell Culture Services, West Lebanon, NH. Splenocytes were isolated from these animals at three days post MAb injection and stained with a phycoerythrin (PE)-conjugated rat anti-mouse CD4 antibody (PharMingen) and an allophycocyanin (APC)-conjugated rat anti-mouse CD8 antibody (PharMingen). CD4⁺ and CD8⁺ cell depletion from the splenocytes were assessed by a FACS Calibur flow cytometer (BD Biosciences, San Jose, CA) and analyzed by using FlowJo analysis software (Tree Star, Inc., San Carlos, CA).

In the depletion-challenge study, VSV-ssLcrV1 vaccinated mice were injected (IP) with 1 mg (500 μ g/day/mouse) of isotype control, or anti-mCD4 and/or anti-mCD8 MAbs, as indicated, one day prior to and on the day of challenge.

2.8. *Y. pestis* challenge

All challenge work with *Y. pestis* was performed in a biosafety cabinet in a BSL3-equipped animal facility at the PHRI, NJ. Mice vaccinated IM were challenged with 10 LD₅₀ (10,000 cfu) of *Y. pestis* (C092 strain) by the IN route in a volume of 20 μ L/nostril, following anesthetization with Ketamine and Xylazine (40–80 mg/kg and 5 mg/kg of body weight, respectively). Liquid inocula (20 μ L) were delivered to the nares of the mice 5.0 μ L at a time using a pipettman with an aerosol barrier tip. Mice were challenged at the indicated times post-inoculation and were observed daily for up to 14 days. Any animal exhibiting extreme lethargy or distress during this period was euthanized.

3. Results

3.1. Construction and characterization of a recombinant VSV expressing a secreted form of the *Yersinia pestis* LcrV protein

To generate a recombinant VSV that expressed a secreted form of the *Y. pestis* LcrV protein, a DNA sequence encoding signal sequence (ss) for secretion was added upstream of the LcrV ORF and inserted into the plasmid vector pVSV1XN [51]. This allowed the recovery of a virus, VSV-ssLcrV1 (Fig. 1A), expressing a secreted form of LcrV from the first position of the VSV genome, the position that maximizes mRNA and protein expression. VSV transcription occurs sequentially starting from the 3'-end, with transcription being attenuated by about 30% at each subsequent gene junction [53]. Appropriate VSV-specific transcription start, stop, and polyadenylation signals flanking the ssLcrV-coding region were incorporated as indicated (Fig. 1A). The recovered virus grew to titers nearly equivalent to wild-type VSV and had a normal plaque size.

To verify expression of the secreted LcrV protein and to examine its localization in cells, we infected cells with VSV-ssLcrV1, VSV-LcrV1, or recombinant wild-type (rwt)-VSV and then performed indirect immunofluorescence microscopy using two monoclonal antibodies (MAbs) to LcrV protein. As shown in Fig. 1B, the protein encoded by VSV-ssLcrV1 was present in a reticular network throughout the cytoplasm, the pattern expected for a secreted protein in the endoplasmic reticulum. In contrast, the LcrV protein encoded by VSV-LcrV1 showed a more diffuse cytoplasmic localization. Control cells infected with rwt-VSV showed no staining for LcrV protein. Differential interference contrast (DIC) images are included to show the presence of cells in all fields.

The expression of the secreted LcrV protein was also verified by metabolically labeling VSV-ssLcrV1 infected BHK-21 cells with [³⁵S]-methionine, followed by analysis of lysates and supernatants from the infected cells by SDS-PAGE and autoradiography (Fig. 2). Because VSV shuts off host protein synthesis, the cell lysates from rVSV- infected cells show mainly the viral encoded proteins L, G, N, P and M (Fig. 2). High-level expression of a protein with the expected mobility of LcrV (~37kDa) was identified both in cell lysates and in supernatants from cells infected with VSV-ssLcrV1 (Fig. 2, lanes 3 and 7), confirming the secretion of the ssLcrV protein. Control cells infected with VSV-LcrV1 [24], expressed the protein only in the cell lysates and not in the medium (Fig. 2, lanes 4 and 8). The LcrV band was absent in lysates and supernatants from mock infected cells or cells infected with rwt-VSV (Fig. 2, lanes 1, 2, 5 and 6).

3.2. Intramuscular vaccination with VSV-ssLcrV1 elicits improved antibody response to LcrV with increasing immunizing dose

Because our earlier work showed that, a single intranasal (IN) immunization (10^6 pfu) with VSV-LcrV1 generated low-level and variable protection (10%) to *Y. pestis* challenge [24], we generated the vector expressing secreted LcrV (VSV-ssLcrV1) and evaluated the antibody response generated by this vector. A single IN vaccination with a 10^6 pfu of VSV-ssLcrV1 generated a modestly improved protection against intranasal *Y. pestis* challenge when compared to VSV-LcrV1 (not shown). To determine if better protection could be obtained using a higher immunizing dose, we initiated a dose response study. Previous studies with VSV vectors have shown little effect of virus dose when given intranasally [37], but substantial effects of dose when given by the intramuscular (IM) route (S. Kapadia and J.K. Rose, unpublished results). We therefore tested the effect of virus dose on the antibody response to VSV-ssLcrV1 given IM.

In an initial study, five groups of mice (five mice per group) were immunized IM with escalating doses (10^5 , 10^6 , 10^7 , 10^8 , or 10^9 pfu) of VSV-ssLcrV1. A group of three mice received the control virus VSV-eGFP1 by the same route. Blood was collected from these mice at 28 and 53 days post immunization (dpi) and anti-LcrV titers in pooled sera were determined using ELISA. As shown in Fig. 3, each escalating IM dose of VSV-ssLcrV1 ranging from 10^5 – 10^9 pfu elicited increasing anti-LcrV titers that improved with time. The increase was more prominent in the two higher dose groups (10^8 and 10^9 pfu), with over two-fold increase in anti-LcrV titers in the 10^9 pfu group, relative to the 10^8 pfu group.

In order to determine if the higher doses of VSV-ssLcrV1 might be potentially pathogenic in mice, we also weighed the mice daily for two weeks following VSV immunizations (not shown). The amount and kinetics of weight loss in mice are a sensitive measure of viral pathogenesis [30]. While weight loss was negligible in mice immunized with 10^5 and 10^6 pfu, similar to that receiving the control virus VSV-eGFP1, mice in 10^7 and 10^8 pfu dose groups showed a transient weight loss (up to 5% by day 1 post immunization) and began to recover by day 2. Mice in the 10^9 pfu dose group showed the greatest weight loss (up to 10% by day 2), but these and all other mice recovered to their pre-immunization body weights by day four.

3.3. A single dose of VSV-ssLcrV1 confers high-level protection against *Y. pestis* challenge

Because the dose response study showed that we could achieve very high anti-LcrV titers with a single high dose of VSV-ssLcrV1, we compared antibody responses and degree of protection from *Y. pestis* challenge conferred by increasing doses of VSV-ssLcrV1 and VSV-LcrV1, the vector that we had used earlier [24]. The immunization timeline for this challenge study is represented in Fig. 4A. Three groups of eight mice each were vaccinated (IM) with VSV-ssLcrV1, at doses of 10^7 , 10^8 and 10^9 pfu. In parallel, three additional groups received the same doses of VSV-LcrV1. Eight control mice were immunized (IM) with VSV-eGFP1. Blood was collected at indicated time-points (Fig. 4A) and anti-LcrV titers were determined using pooled sera from each immunization group. Increasing doses of either virus led to increased antibody titers. In addition, the titers induced by VSV-ssLcrV1 were generally higher than those induced by VSV-LcrV1 (Fig. 4B).

Analysis of IgG subclass binding LcrV protein (10^9 pfu dose) showed that VSV-ssLcrV1 vaccination induced an LcrV-specific IgG2a response which was about 3-fold higher than the IgG1 response (Fig. 4C). In contrast, the VSV-LcrV1 vaccination induced similar levels of LcrV-specific IgG1 and IgG2a (Fig. 4C).

To measure the efficacy of these single dose recombinant VSV vaccines, animals from these immunization groups were subjected to pulmonary challenge (IN) with 10 LD₅₀ *Y. pestis* (CO92 strain) at about 3 months post vaccination. In the VSV-ssLcrV1 groups, seven out of eight animals in the 10^9 pfu group survived the challenge (Fig. 5A). Four out of eight animals survived in the 10^8 pfu group, while only one out of eight mice survived in the 10^7 pfu group (Fig. 5A). All mice in the VSV-eGFP1 control group died by day 5 post-challenge. The higher level of protection in the 10^9 pfu group was statistically highly significant [$p = 0.001$, Fisher Exact Test (when compared to control vaccinated animals)]. In comparison, in both 10^9 and 10^8 pfu groups of the VSV-LcrV1 set, only three of eight animals survived the challenge, while none survived in the 10^7 pfu group (Fig. 5B).

The difference in survival between the VSV-ssLcrV1 vaccinated and VSV-LcrV1 vaccinated animals (10^9 dose) approached statistical significance in this first experiment ($p=0.059$). In a second experiment (10 animals per group) using the same dose of each vector and the same challenge, we obtained 100% protection in the VSV-ssLcrV1 vaccinated animals and only 50% protection in the VSV-LcrV1 vaccinated animals. This difference was highly significant ($p = 0.016$).

3.4. Survival did not fully correlate with antibody titer

Earlier passive transfer studies have shown that antibody to LcrV is sufficient to provide protection from disease in the mouse model [21,23]. Although in our studies, increased antibody titers to LcrV generally yielded higher protection, analysis of antibody responses in individual mice often did not correlate well with protection. Comparison of mouse survival profiles to their anti-LcrV titers in the 10^8 pfu dose groups, revealed that there was a large variation in the titers among individual mice and that some mice with low titers survived the *Y. pestis* challenge (Fig. 6).

3.5. Role of CD4⁺T-cells in protection elicited by VSV-ssLcrV1 vaccination

Because LcrV antibody titers did not correlate completely with protection (Fig. 6), we thought that cellular immune responses might contribute to protection against pulmonary plague infection induced by VSV vectors. To examine a potential role of CD4⁺ and CD8⁺ cells in VSV-ssLcrV1 vaccinated mice at the time of challenge we depleted these cells singly or together just before challenge and then determined the extent of protection. A control experiment was performed to evaluate the extent of depletion using mouse specific MAbs. Figure 7 shows representative FACS plots of splenocytes, obtained from mice injected intraperitoneally (IP) with anti-mCD4, anti-CD8 or isotype control MAbs and, stained with appropriately conjugated rat anti-mouse CD4 or CD8 antibodies. Relative to an isotype control MAb treated mouse, anti-mCD4 and/or anti-mCD8 MAb treatment resulted in a nearly complete, specific depletion of CD4⁺ and/or CD8⁺ cells in mice, confirming the efficiency of the depletion (Fig. 7).

Figure 8A gives the plan and timeline for the depletion/challenge study. Four groups of ten mice each (groups A-D) were given 10^9 pfu of VSV-ssLcrV1 by the IM route, while ten control mice received similar doses of VSV-eGFP1. The CD4⁺ and/or CD8⁺ cells were depleted in some of these immunization groups with injection (IP) of MAbs specific for mouse CD4 and/or mouse CD8 as indicated (Fig. 8A), followed by pulmonary *Y. pestis* challenge approximately 3 months post vaccination. Complete protection [$p = 0.03$ (Fisher Exact Test), as compared to VSV-eGFP1 vaccinated mice] was obtained in the VSV-ssLcrV1 vaccinated animals, which received the isotype control MAb (group A) or the anti-mCD8 MAb (group C), as shown in Fig. 8B. In contrast, two groups of vaccinated mice that received the anti-mCD4 MAb, with or without depletion of CD8⁺ cells (groups B and D), elicited significantly reduced [$p=0.02$ (Fisher Exact Test)] levels of protection (70%). Note that greater statistical significance was achieved in the latter case because there were more animals involved in this calculation. Surprisingly, one out of the ten control animals survived the challenge. In our previous studies of mice given this challenge, all control mice ($n=35$) had died by day 4 or 5 post challenge.

4. Discussion

In an earlier study, we showed that an intranasal (IN) prime-boost immunization with two different VSV vectors expressing LcrV elicited high-level protection against IN *Y. pestis* challenge [24]. We report here the development of an rVSV vector (VSV-ssLcrV1), expressing a secreted form of LcrV from the first position of the VSV genome, that when given intramuscularly (IM) as a single dose of 10^9 pfu, induced high-level anti-LcrV antibody titers in mice, and completely protected animals against subsequent lethal pulmonary *Y. pestis* challenge. Compared to the VSV-LcrV1 vector (expressing non-secreted LcrV), that we described earlier [24] the vector expressing secreted LcrV (VSV-ssLcrV1) gave significantly better single-dose protection. Immunization with VSV-ssLcrV1 also elicited higher IgG2a relative to IgG1 titers suggestive of a shift toward a T_H1 type of immune response to the secreted protein.

Although the basis for protection by LcrV-based vaccines is not completely understood, several studies have shown a correlation between high anti-LcrV titers and protection [15,16,25,26]. Additionally, passive transfer of polyclonal LcrV specific antibody, as well as LcrV monoclonal antibody (MAb) treatment was shown to be protective in mouse models of plague infection that LcrV antibody can be sufficient for protection [22,23].

In our study, we observed that the pre-challenge LcrV ELISA titers did not correlate completely with protection because animals having even very low titers were protected in some instances. It was also noted in a non-human primate model of plague infection that some animals possessing high LcrV antibody titers succumbed to *Y. pestis* challenge [54]. Together these results suggested a possible cellular immune component in protection. Recently, immunizations with live replicating *Y. pestis* have also shown the importance of cell-mediated immunity in protection against plague challenge [55,56]. However, LcrV did not appear to be a major inducer of T-cell responses in that study [56].

In our studies, induction of increased IgG2a response to LcrV following immunization with VSV-ssLcrV1 indicated an antigen-specific bias toward a T_H1 cellular immune response. To determine if we could identify a role for cellular immunity in protection, we performed selective depletion of mouse $CD4^+$ and/or $CD8^+$ T-cells at the time of challenge. Mice completely depleted for $CD8^+$ T-cells prior to challenge were 100% protected from disease indicating that $CD8^+$ T-cells are not required for protection. However, depletion of $CD4^+$ T-cells, with or without depletion of $CD8^+$ T cells, significantly reduced the level of protection. This result suggests that $CD4^+$ T-cells might be involved in protection, although effects on other $CD4^+$ cells cannot be ruled out. Effects on antibody production are unlikely because the depletion is performed just prior to challenge. Consistent with our findings, $CD4^+$ T-cell epitopes have been reported for the LcrV protein in Balb/c mice [57], while $CD8^+$ T-cell epitopes have not been defined.

What role might the $CD4^+$ T-cells be playing in protection against *Y. pestis*? Vaccine induced $CD4^+$ T-cells are known to secrete phagocyte activating type 1 cytokines like interferon gamma ($IFN-\gamma$) and tumor necrosis factor alpha ($TNF-\alpha$), which contribute to intracellular antimicrobial defense by activating macrophages [58,59]. However, although intracellular niches of *Y. pestis* exist, pulmonary plague infection is considered predominantly extracellular. Some recent studies suggest that both $CD4^+$ T cells and $IFN-\gamma$ play critical roles in combating extracellular bacterial infections [60,61]. Other studies imply important roles for $IFN-\gamma$ and $TNF-\alpha$ in protection against lethal *Y. pestis* infections [62,63]. Thus depletion of vaccine primed $CD4^+$ T-cells leading to abrogation of the cytokine secreting activity, might be reflected in the reduced level of protection that we have observed. In addition, the role of cytotoxic $CD4^+$ T-cells in eliminating infected cells also cannot be ruled out.

Why addition of an amino terminal signal sequence for secretion of the LcrV protein enhances protection is unclear, but this was also noted previously for multi-dose DNA vaccine expressing secreted LcrV [26]. The authors of that study suggested that aggregation of the secreted protein might be involved in greater induction of antibody. The inclusion of the signal sequence, which we show here causes efficient secretion of LcrV, could also be responsible for enhancing $CD4^+$ T-cell responses if the secreted protein were more efficiently directed into the MHC class II presentation pathway.

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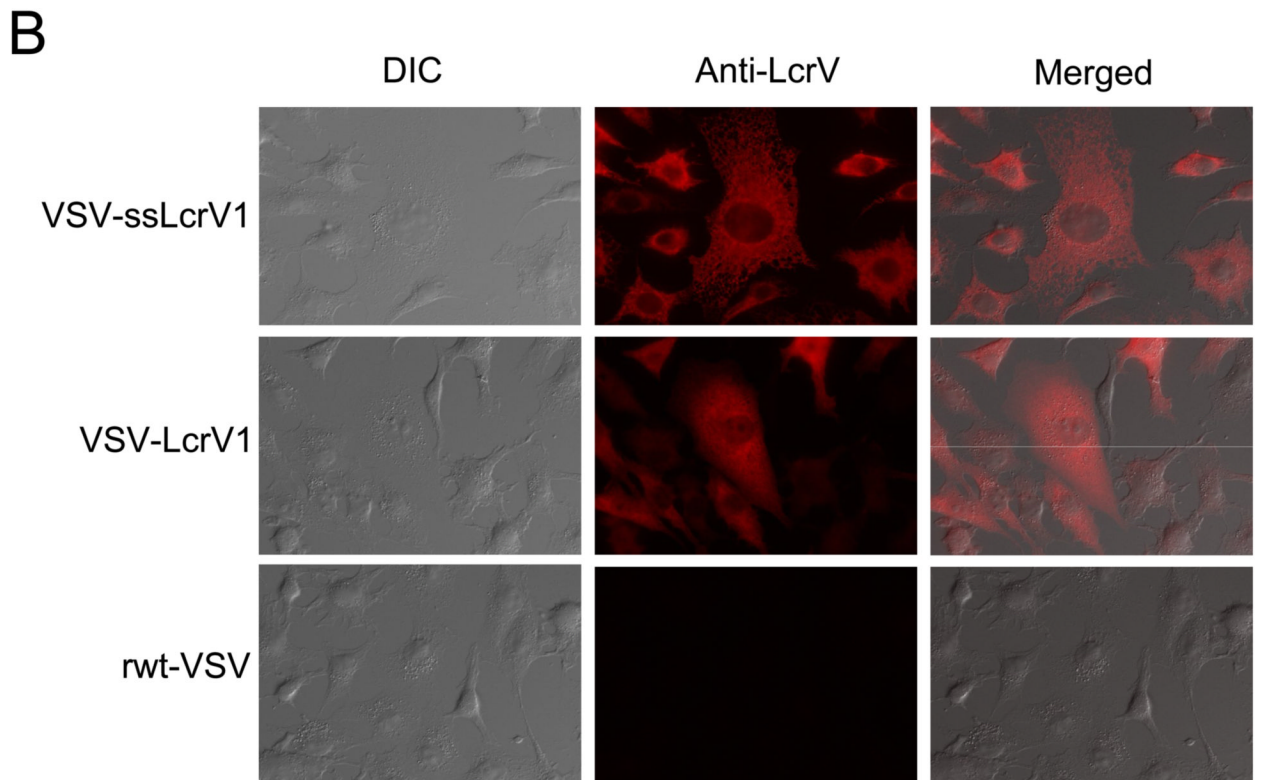
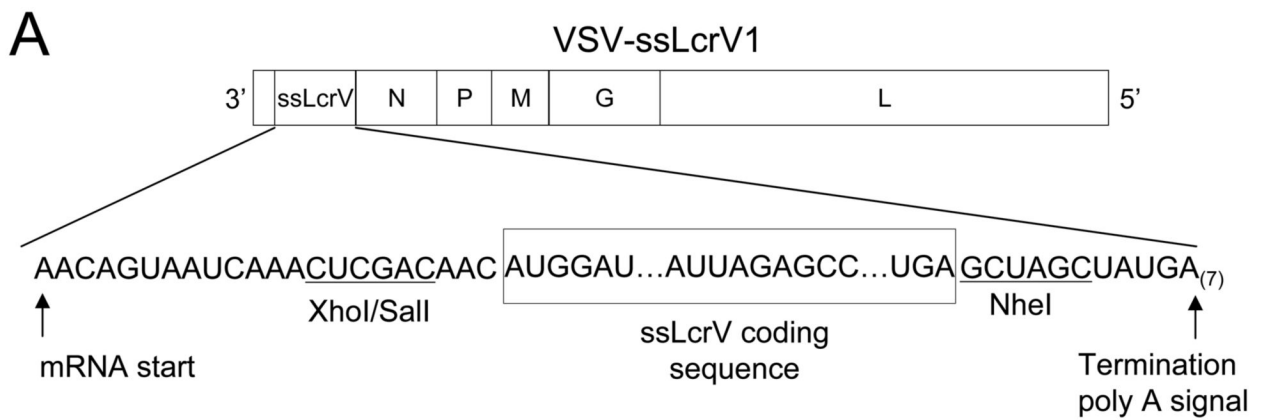


Figure 1. Recombinant VSV expressing the secreted form of *Yersinia pestis lcrV* gene (A) Schematic representation of the recombinant VSV with ssLcrV inserted upstream of the N gene (VSV-ssLcrV1) used in this study, showing the gene order in 3' to 5' direction on the negative-strand genome. (B) Indirect immunofluorescence microscopy of BHK-21 cells infected with the indicated viruses. Cells were fixed at 3 hrs post infection, permeabilized and stained using anti-LcrV monoclonal antibodies (MAbs), followed by an AlexaFlour-594 secondary antibody. The differential interference contrast (DIC) images are shown in the left column, the fluorescence images are shown in the middle column and the merged images are shown in the right column.

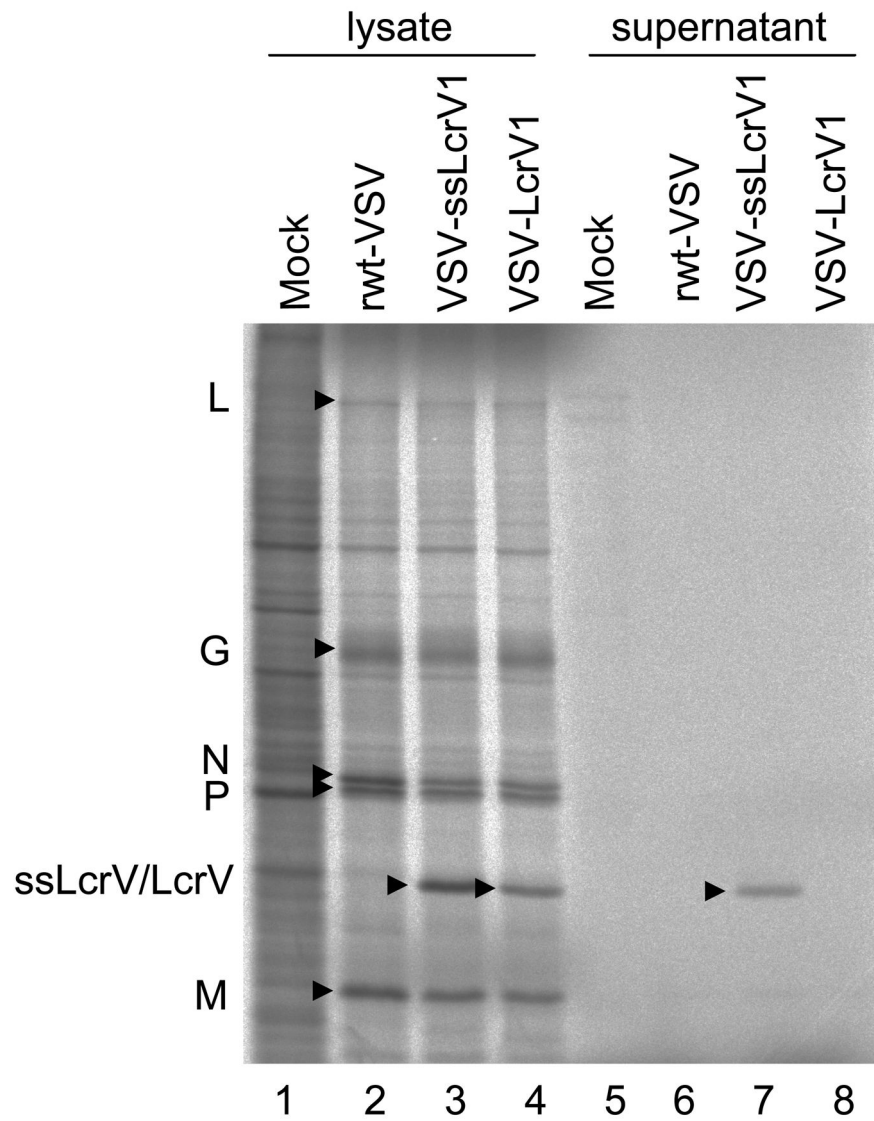


Figure 2. Metabolic labeling of proteins expressed from recombinant VSV vectors. BHK-21 cells, were infected with the indicated viruses or mock infected and then metabolically labeled with [³⁵S]-methionine. Cell lysates and cell supernatants were fractionated by 10% SDS-PAGE and the gel image collected on a phosphorimager. Positions of VSV encoded proteins are indicated on the left side of the gel image.

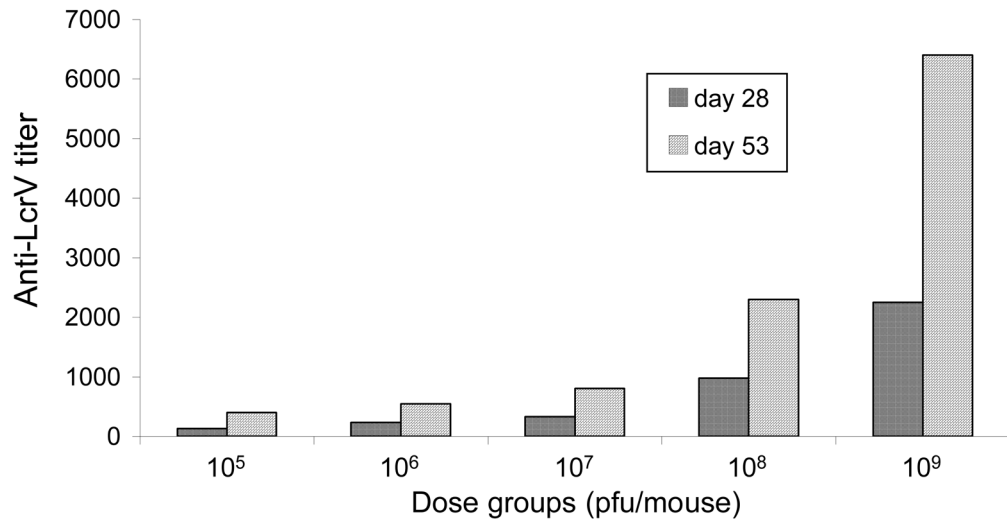


Figure 3. Dose-response of anti-LcrV serum titers in VSV-ssLcrV1 immunized mice. Mice in five different groups were immunized intramuscularly (IM) on day 0 with 10⁵, 10⁶, 10⁷, 10⁸ and 10⁹ pfu of VSV-ssLcrV1. Blood was collected, by retro-orbital bleeding, on days 28 and 53 post inoculation. Serum samples were pooled and total IgG titers, specific to LcrV, were determined at the indicated days using ELISA.

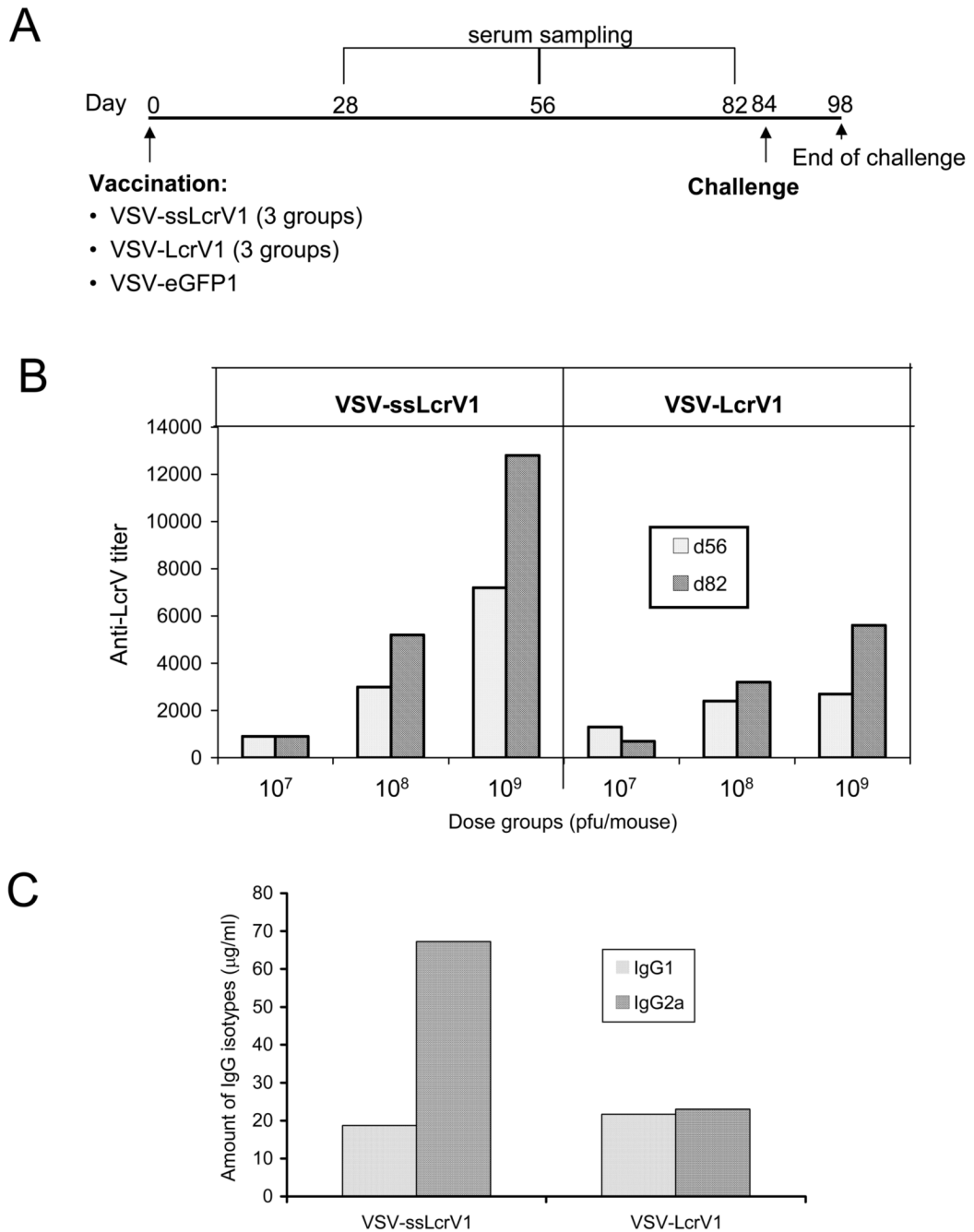


Figure 4.

Comparison of humoral immune responses of mice to VSV-ssLcrV1 and VSV-LcrV1 vaccinations. (A) Timeline for dose response vaccination-challenge study. Mice were vaccinated IM with 10⁷, 10⁸ and 10⁹ pfu doses of rVSVs encoding either ssLcrV or LcrV and challenged IN with *Y. pestis* (CO92 strain). VSV-eGFP1 immunization (10⁷ pfu) was used as control. (B) Anti-LcrV titer profile at 56 and 82 days post immunization, in mice immunized with 10⁷, 10⁸ and 10⁹ pfu doses of either VSV-ssLcrV1 (left panel) or VSV-LcrV1 (right panel). (C) Amount of anti-LcrV specific IgG1 and IgG2a isotypes in VSV-ssLcrV1 (left) of VSV-LcrV1 (right) vaccinated mice in the 10⁹ pfu dose group. Serum antibodies that bound LcrV were quantitated using secondary IgG1- and IgG2-specific antibodies. Concentrations

of mouse IgG1 or IgG2a were then calculated from standard curves run in parallel using a range of concentrations of purified mouse IgG1 or IgG2a.

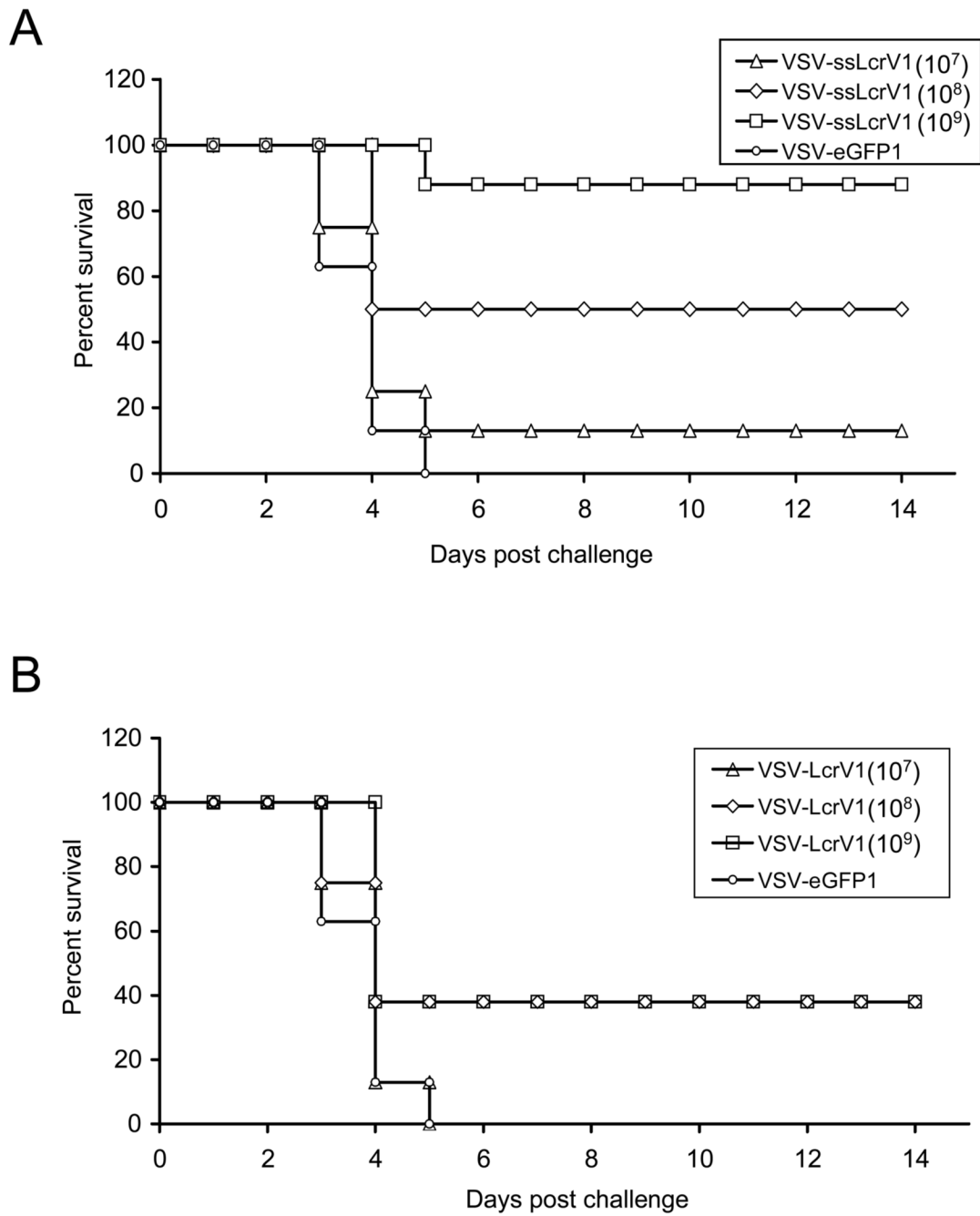


Figure 5. Protective efficacy of single-dose VSV-ssLcrV or VSV-LcrV vaccine in a plague challenge assay. (A) Survival profile of VSV-ssLcrV1 vaccinated mice following lethal plague challenge. Mice were immunized (IM) with VSV-ssLcrV1 at three different doses - 10⁷ pfu (open triangle), 10⁸ pfu (open diamond) or 10⁹ pfu (open square). The control mice received VSV-eGFP1 (open circle). All animals were challenged at 3 months post vaccination with 10LD₅₀ *Y. pestis* (CO92) and observed for 14 days. (B) Similar to A, except that animals were vaccinated with three different doses of VSV-LcrV1 - 10⁷ pfu (open triangle), 10⁸ pfu (open diamond) or 10⁹ pfu (open square).



Figure 6. Correlation of pre-challenge anti-LcrV titers and mouse survival. Mice were vaccinated with 10^8 pfu of either VSV-ssLcrV1 or VSV-LcrV1 and challenged at day 84 post vaccination with 10 LD₅₀ (10,000 cfu) of *Y. pestis* (CO92 strain). Anti-LcrV serum titers in individual mice at day 82 post vaccination are plotted for those mice that survived challenge (alive, filled triangle) or succumbed to challenge (dead, open inverted triangle). The horizontal bars represent the geometric means for the two groups.

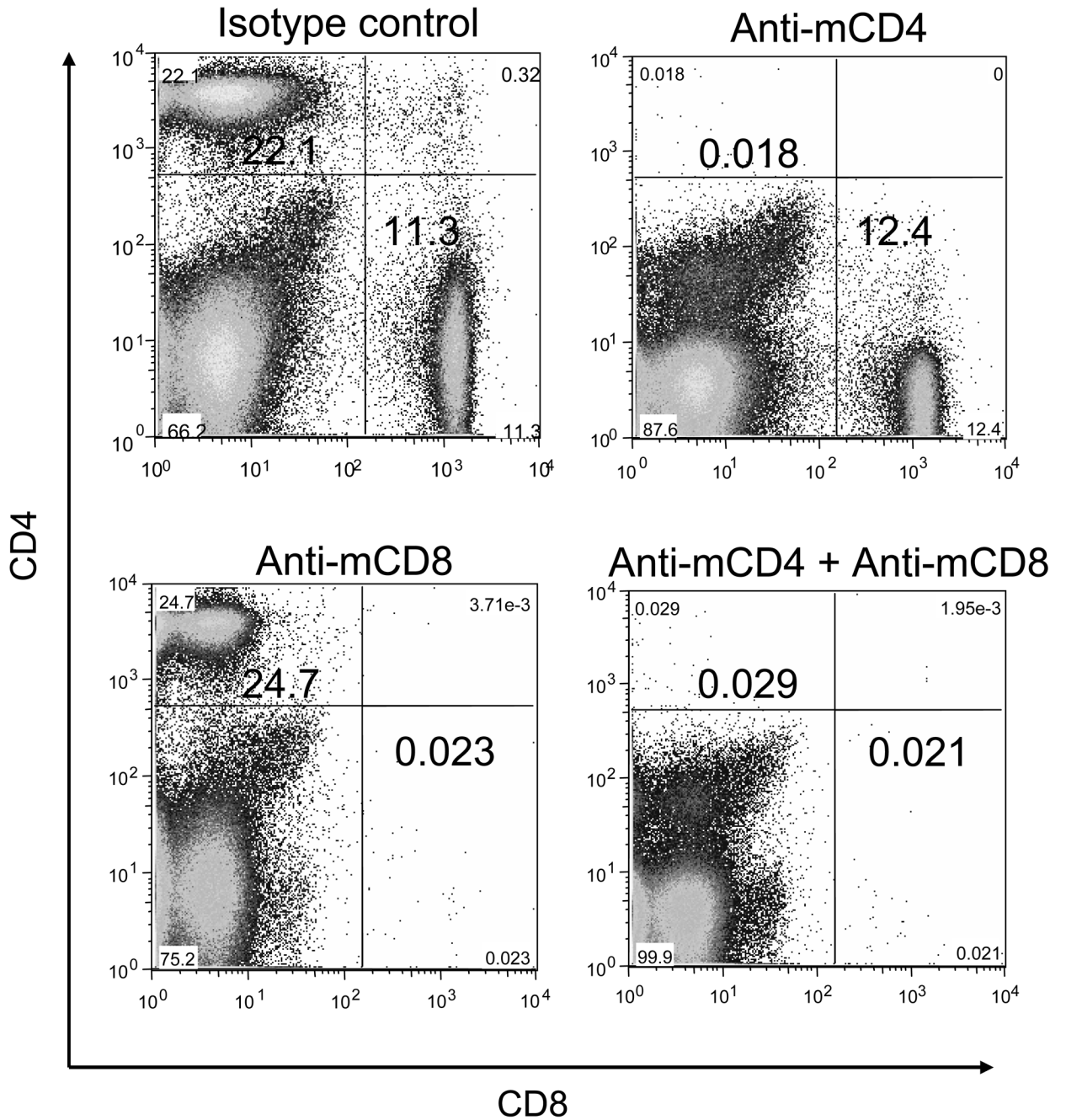
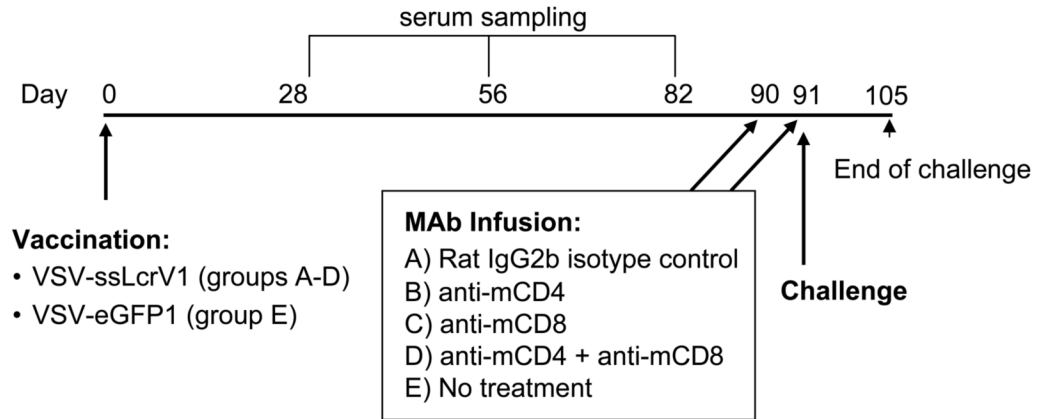


Figure 7. Depletion of CD4⁺ and CD8⁺ T-cells in mice using mouse specific MAbs. Representative FACS plots are shown for splenocytes obtained from mice injected intraperitoneally (IP) with anti-mouse isotype control (top left), anti-mCD4 (top right), anti-mCD8 (bottom left) or combined anti-mCD4 + anti-mCD8 (bottom right) MAbs and stained with appropriately conjugated rat anti-mouse CD4 and CD8 antibodies on day 3 post injection.

A



B

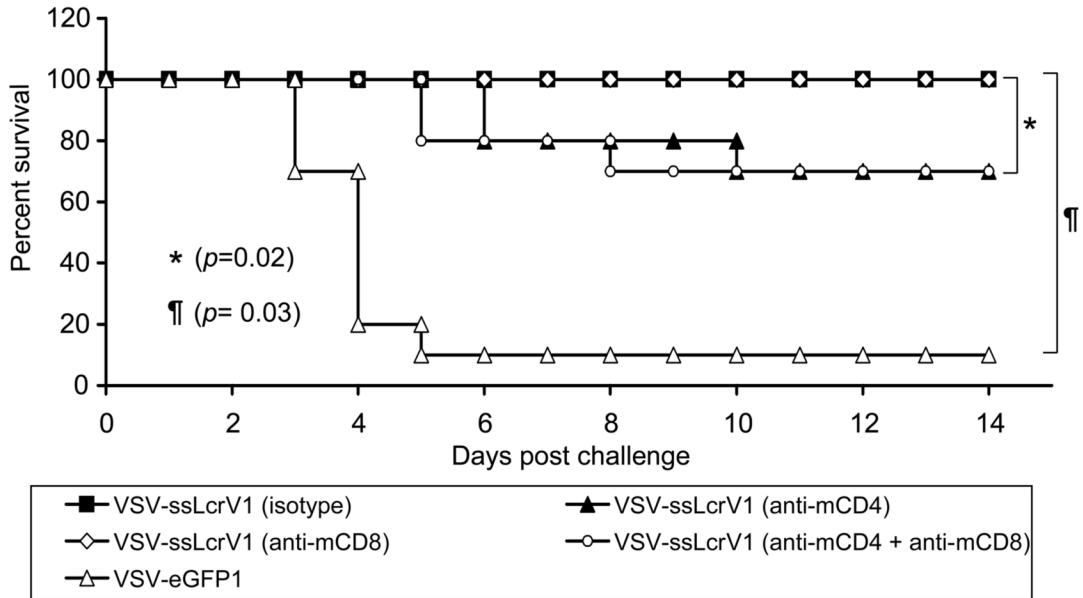


Figure 8.

CD4⁺ cells are required at time of challenge for complete protection against plague challenge. (A) Timeline for the depletion-challenge study. Mice in four groups (A-D) were immunized IM with 10⁹ pfu of VSV-ssLcrV1. Serum sampling was done on day 28, 56 and 82 post immunization. Mice were injected with CD4⁺/CD8⁺ depleting MAbs (IP) on days 90 and 91 post immunization followed by challenge on day 91 with 10 LD₅₀ (10,000 cfu) of *Y. pestis* (CO92 strain). VSV-eGFP1 vaccinated mice were used as the control group. Rat IgG2b isotype MAb was used as a control. (B) Survival profile for mice in the depletion-challenge experiment. VSV-ssLcrV1 vaccinated mice were injected with isotype-matched control (filled square),

anti-mCD4 (filled triangle), anti-mCD8 (open diamond) or mCD4+mCD8 (open circle) MAbs. Control animals receiving VSV-eGFP1 (open triangle) were not treated with MAbs.