

A Viable Recombinant Rhabdovirus Lacking Its Glycoprotein Gene and Expressing Influenza Virus Hemagglutinin and Neuraminidase Is a Potent Influenza Vaccine

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ABSTRACT

The emergence of novel influenza viruses that cause devastating human disease is an ongoing threat and serves as an impetus for the continued development of novel approaches to influenza vaccines. Influenza vaccine development has traditionally focused on producing humoral and/or cell-mediated immunity, often against the viral surface glycoproteins hemagglutinin (HA) and neuraminidase (NA). Here, we describe a new vaccine candidate that utilizes a replication-defective vesicular stomatitis virus (VSV) vector backbone that lacks the native G surface glycoprotein gene (VSVΔG). The expression of the H5 HA of an H5N1 highly pathogenic avian influenza virus (HPAIV), A/Vietnam/1203/04 (VN1203), and the NA of the mouse-adapted H1N1 influenza virus A/Puerto Rico/8/34 (PR8) in the VSVΔG vector restored the ability of the recombinant virus to replicate in cell culture, without the requirement for the addition of trypsin. We show here that this recombinant virus vaccine candidate was non-pathogenic in mice when given by either the intramuscular or intranasal route of immunization and that the *in vivo* replication of VSVΔG-H5N1 is profoundly attenuated. This recombinant virus also provided protection against lethal H5N1 infection after a single dose. This novel approach to vaccination against HPAIVs may be widely applicable to other emerging strains of influenza virus.

IMPORTANCE

Preparation for a potentially catastrophic influenza pandemic requires novel influenza vaccines that are safe, can be produced and administered quickly, and are effective, both soon after administration and for a long duration. We have created a new influenza vaccine that utilizes an attenuated vesicular stomatitis virus (VSV) vector, to deliver and express influenza virus proteins against which vaccinated animals develop potent antibody responses. The influenza virus hemagglutinin and neuraminidase proteins, expressed on the surface of VSV particles, allowed this vaccine to grow in cell culture and induced a potent antibody response in mice that was effective against infection with a lethal influenza virus. The mice showed no adverse reactions to the vaccine, and they were protected against an otherwise lethal influenza infection after only 14 days postvaccination and after as many as 140 days postvaccination. The ability to rapidly produce this safe and effective vaccine in cell culture is additionally advantageous.

With the ongoing public health threat of emerging strains of influenza virus and the precedent for potentially devastating worldwide pandemics, novel approaches to influenza vaccination remain a priority. The influenza virus surface glycoproteins hemagglutinin (HA) and neuraminidase (NA) are key mediators of the viral infectious cycle and obvious proteins of interest for novel vaccines. The majority of recent vaccine candidates focus predominantly on the immunogenic properties of the influenza virus HA, and sometimes NA, glycoprotein, seeking to utilize vector systems (1–3), adjuvants (4–6), alternative routes of immunization (7, 8), and novel antigen presentation and delivery systems (9–12) to increase humoral and/or cell-mediated immunity against these and other influenza virus antigens. Creating vaccines that also exploit the functional properties of the influenza virus glycoproteins represents a unique approach.

Influenza virus propagation in tissue culture typically requires the addition of exogenous trypsin to cleave the HA0 precursor, commonly at a basic amino acid between HA1 and HA2. This cleavage is essential for virus attachment and entry into host cells (13). Highly pathogenic avian influenza viruses (HPAIVs) have been shown to contain polybasic amino acid cleavage sites in HA

that can be promiscuously activated by ubiquitous cellular proteases, leading to severe disease and also allowing for trypsin-independent growth in tissue culture (14–17). These influenza viruses have caused devastating outbreaks among poultry and wild birds and have sporadically caused cases of human disease (18–21). HPAIVs were originally found in areas of China and Southeast Asia, where they still circulate today, although their geographic distribution has spread widely (21, 22).

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Our laboratory has developed vesicular stomatitis virus (VSV) as a vector for a wide variety of experimental vaccines (23–32). Due to its lack of pathogenicity in humans, potent induction of immune responses, wide range of cellular tropism, and exuberant growth properties, VSV is an ideal vaccine vector. Although the results of the first phase I trial utilizing an attenuated VSV vector as an HIV-1 vaccine candidate have not yet been published, preliminary safety and immunogenicity results suggest that this vaccine does not cause significant adverse events in humans (HVTN 090 study [<http://clinicaltrials.gov/ct2/show/NCT01438606>]).

VSV relies on the G surface glycoprotein for infectivity. Replication-deficient, pseudotyped VSVΔG vaccine vectors have been shown to be safe and effective in animal models (31, 33–35). While VSVΔG strains pseudotyped with VSV G or other viral glycoproteins are capable of only a single round of infection, our laboratory and others have shown that substituting other viral surface glycoprotein genes for VSV G in recombinant viruses can restore replication competence to VSVΔG (23, 36–39). We have termed these recombinants “surrogate viruses” to distinguish them from VSV pseudotypes that are made by complementation of VSVΔG with other viral glycoproteins (38). These recombinant, surrogate viruses can serve as potent vaccines while lacking pathogenicity. In previous studies, we generated VSVΔG recombinants expressing influenza virus HA and NA genes from a mouse-adapted influenza virus (WSN), but these viruses did not propagate without VSV G complementation, even though substantial quantities of HA and NA were present in the viral membrane (our unpublished data).

In the current study, we have created a novel vaccine against HPAIVs utilizing a hybrid VSVΔG vector expressing the HA of the H5N1 HPAIV A/Vietnam/1203/04 (VN1203) and the NA of a mouse-adapted H1N1 influenza virus, A/Puerto Rico/8/34 (PR8). This virus propagates in BHK-21 cells without any requirement for the addition of exogenous trypsin. This surrogate virus is non-pathogenic in mice yet protects mice against lethal challenge with an H5N1 influenza challenge virus (HALo, a 6:2 reassortant influenza virus containing the HA and NA segments of VN1203 and internal gene segments from PR8, with the removal of the polybasic cleavage site in HA [40]).

MATERIALS AND METHODS

Plasmid constructions. To generate pVSVΔG-H5N1, the VSV G gene was first removed from full-length plasmid pVSV-H5-HA(VN1203) (27) by MluI-XhoI digestion. The PR8 NA gene was amplified by PCR from pCAGGS-NA(PR8) using the forward primer GTACGTACacgctactA TGAATCCAAATCAGAAAAT and the reverse primer GATCGATCctcgagCGTGATATCTGTTAGTTTTTTCATACTACTTGTCAATGC TGAATGG (lowercase letters represent restriction sites), containing a VSV transcription stop-start sequence. This PCR product was purified, digested with MluI-XhoI, and cloned into the VSVΔG vector described above. Plasmid pCAGGS-HA(VN1203) was made by digesting pVSV-H5-HA(VN1203) with XhoI-NheI to release the H5 HA gene, which was then ligated into pCAGGS digested with XhoI-NheI.

Metabolic labeling of cells infected with recombinant viruses and SDS-PAGE of cell lysates. BHK-21 cells were infected with recombinant viruses at a multiplicity of infection (MOI) of 10. At 6 h postinfection (p.i.), cells were washed with methionine-free Dulbecco’s modified Eagle’s medium (DMEM), followed by labeling with 100 μCi of [³⁵S]methionine in 0.5 ml of methionine-free DMEM for 1 h at 37°C. Cells were washed with phosphate-buffered saline (PBS) and lysed with detergent lysis buffer (1% Nonidet P-40, 0.4% deoxycholate, 50 mM Tris-HCl [pH 8.0], 62.5 mM EDTA) on ice for 5 min. Lysates were subjected to SDS-

PAGE on a 4 to 12% Bis-Tris NuPAGE gel (Invitrogen, CA). Protein bands on the dried gel were scanned by using a Fujifilm BAS 1800 imaging system.

Antisera and antibodies. The anti-VSV G monoclonal antibodies I1 and I14 (41) were used to detect VSV G expression. Sheep antiserum to recombinant HA (rHA) [subtype H5 influenza A/HK/156(483)/97 (H5N1) virus] was obtained from BEI Resources. Polyclonal anti-VSV serum was used to detect VSV proteins in Western blots. Antisera to PR8 NA was obtained from BALB/c mice that had recovered from an infection with PR8.

Indirect immunofluorescence microscopy. BHK-21 cells on coverslips were infected with recombinant wild-type (wt) VSV, VSV-H5, or VSVΔG-H5N1 for 6 h. Cells were then washed twice with PBS and fixed with 3% paraformaldehyde for 30 min. Cells were again washed with PBS-glycine (10 mM) and incubated with a 1:200 dilution of VSV G monoclonal antibodies (I1/I14), a 1:200 dilution of sheep anti-H5N1, or a 1:100 dilution of mouse anti-PR8 NA. The cells were then washed in PBS-glycine and incubated with a 1:500 dilution of goat anti-mouse Alexa Fluor 488 IgG or donkey anti-sheep Alexa Fluor 488 IgG (Molecular Probes, Eugene, OR). Cells were washed twice in PBS-glycine, mounted onto slides by using Prolong Gold antifade reagent with 4’,6-diamidino-2-phenylindole (DAPI) (Molecular Probes, Eugene, OR), and imaged with a Nikon Eclipse 80i fluorescence microscope using a 40× objective.

Recombinant virus recovery. Recombinant VSV (rVSV) was recovered from pVSVΔG-H5N1 as described previously (42, 43). In brief, BHK-21 cells were infected with vTF-7.3 (44) at an MOI of 10. The cells were transfected with pVSVΔG-H5N1 together with support plasmids pBS-N, pBS-P, pBS-L, and pBS-G, encoding VSV proteins. The cell supernatants were collected at 48 h and passaged onto BHK-21 cells that were transfected with pCAGGS-G (45) 1 day earlier. This supernatant containing the virus was collected after 48 h. The virus was plaque purified without VSV G complementation on BHK-21 cells and further passaged on BHK-21 cells to generate a non-G-complemented VSVΔG-H5N1 stock. The stock was serially diluted and plaqued on BHK-21 cells. Plaques were stained with crystal violet after 48 h.

Virus neutralization assays. To measure neutralizing antibodies (NAbs) to H5N1 in mice, serum samples were serially diluted with PBS in a 96-well plate and mixed with ~100 infectious particles of wt VSV or VSVΔG-H5N1 to a final volume of 100 μl. The mixture was incubated at 37°C for 1 h, and 100 μl of BHK-21 cells (5 × 10³ cells) was added to each well. The cells were incubated at 37°C for 48 to 72 h. The neutralization titer was defined as the reciprocal of the highest serum dilution at which duplicate wells showed complete neutralization of the virus.

Neutralizing antibody titers were additionally determined by a micro-neutralization assay, as described previously (46). Briefly, serial 2-fold dilutions of heat-inactivated serum were prepared, beginning with a 1:10 dilution. Equal volumes of serum and virus were combined, incubated at room temperature for 1 h, and added to MDCK cells. The residual infectivity of each dilution of the virus-serum mixture was determined in quadruplicate. The neutralizing titer was defined as the reciprocal of the highest dilution of serum that completely neutralized the infectivity of 100 50% tissue culture infective doses (TCID₅₀) of the appropriate wt H5N1 virus for MDCK cells. Infectivity was identified by the presence of a cytopathic effect (CPE) on day 4.

Endpoint ELISA titers. Flat-Bottom Immuno Nonsterile 4 HBX 96-well plates (Thermo Scientific) were coated with 50 μl of recombinant protein diluted in enzyme-linked immunosorbent assay (ELISA) coating buffer (pH 9.4) at a concentration of 2 μg/ml per well and refrigerated at 4°C overnight. Coating buffer was discarded, and wells were blocked with 100 μl of blocking solution (PBS containing 0.1% Tween 20 [T-PBS], 3% goat serum [Gibco], and 0.5% milk powder) for 1 h at room temperature. Another 50 μl of blocking solution was added to the first column of the wells, as was 1.5 μl of mouse serum (starting concentration of 1:100). The samples were 3-fold serially diluted and incubated at room temperature for 2 h. The plates were washed 6 times with T-PBS, and 50 μl of blocking

solution containing anti-mouse IgG (Fab specific)-peroxidase antibody (Sigma) at a concentration of 1:3,000 was added. After 1 h of incubation at room temperature, the plates were washed 6 times with T-PBS and developed with 100 μ l of SigmaFast OPD (Sigma) per well. The developing process was stopped after 10 min with 3 M hydrochloric acid (HCl), and the absorbance at 490 nm was read with a Synergy H1 hybrid multimode microplate reader (BioTek).

The average absorbance plus 3 times the standard deviations of all blanks below an absorbance of 0.065 was calculated and used as the cutoff for endpoint titer analysis.

Hemagglutination inhibition assays. To remove unspecific serum reactivity, mouse sera were trypsin-heat-periodate treated. A one-half volume of L-1-tosylamido-2-phenylethyl chloromethyl ketone (TPCK)-trypsin at a concentration of 8 mg/ml was added to 1 volume of serum, and the mixture was incubated at 56°C for 30 min and then cooled to room temperature. Three volumes of 0.011 M metapotassium periodate (KIO_4) were added, and sera were incubated at room temperature for 15 min. Three volumes of 1% glycerol saline were added, and sera were incubated for another 15 min at room temperature. Two and a half volumes of 85% PBS were added for a final serum dilution of 1:10.

Influenza viruses were prepared at a concentration of 8 hemagglutination units (HAUs) per 50 μ l, and chicken red blood cells (RBCs) were diluted in PBS to a concentration of 0.5%.

Fifty microliters of trypsin-heat-periodate-treated serum was added to the first well of a 96-well V-bottom plate and 2-fold serially diluted in PBS to a final volume of 25 μ l per well. Twenty-five microliters of diluted virus was added to each well, and the contents were mixed and incubated at room temperature for 30 min. Fifty microliters of RBCs was added to each well, and the contents were mixed. After incubation at 4°C for 45 min, the hemagglutination inhibition (HAI) titers were determined. Back titrations of the viruses were performed to confirm a concentration of 8 HAUs/50 μ l.

Western blots. Whole-cell extracts were prepared from BHK-21 cells infected with VSV G, VSV-H5, and VSV Δ G-H5N1. Samples were analyzed by SDS-PAGE on a 4 to 12% Bis-Tris NuPAGE gel (Invitrogen), and Western blot analysis was performed by using antibodies specific for H5 HA or VSV (Indiana serotype). The anti-H5 blot was stripped and re-probed with anti-VSV antibody.

Animal experiments. Six- to eight-week-old female BALB/c mice were obtained from Charles River Laboratories (Wilmington, MA) and housed for 1 week prior to immunization. Animals were housed under biosafety level 2 (BSL-2) conditions in microisolator cages. Viruses were diluted in serum-free DMEM for immunizations. Intranasal (i.n.) inoculations were performed with either VSV Δ G-H5N1 or a control virus (VSV-EGFP [47]) administered in a volume of 25 μ l to animals that were lightly anesthetized with 20% (vol/vol) isoflurane (Baxter) diluted in propylene glycol. For intramuscular (i.m.) immunizations, the VSV Δ G-H5N1 virus, or a control virus [VSV Δ G-G(NiV) (VSV Δ G expressing the G glycoprotein of Nipah virus) (37)], was injected in a volume of 50 μ l into the left hind leg muscle. Blood was collected from the retro-orbital sinus for neutralization assays.

Influenza virus challenges were performed with the low-pathogenicity H5N1 HALO influenza virus (6:2 reassortant influenza virus containing internal gene segments from A/Puerto Rico/8/34 and HA and NA from A/Vietnam/1203/04 with the removal of the polybasic cleavage site in HA [40]). Mice were anesthetized with 20% isoflurane (Baxter) as described above, and the challenge virus was administered in a volume of 50 μ l. The Yale University and Icahn School of Medicine at Mount Sinai Institutional Animal Care and Use Committees approved all immunization and challenge experiments (Yale University protocol permit number 2012-07680), and experiments were done in a biosafety level 2 animal facility in accordance with the regulations of Yale University.

In vivo recovery of VSV Δ G-H5N1. Groups of BALB/c mice were immunized with either VSV Δ G-H5N1 or a control virus (VSV-cH5/1 [1]) by both the i.m. and i.n. routes. Vaccination with VSV Δ G-H5N1 was

done with a 100-fold-higher dose than that used for vaccination with VSV-cH5/1 (2×10^7 PFU of VSV Δ G-H5N1 versus 2×10^5 PFU of VSV-cH5/1). Mice were euthanized either 24 h or 96 h after vaccination, and tissues (lung, spleen, draining lymph nodes, and leg muscle) were isolated and frozen. Homogenates were prepared in 2.5% DMEM, as previously described (48), and frozen at -80°C prior to inoculation of cell monolayers. BHK-21 cells that were plated onto 24-well plates the day prior were then exposed to 20 μ l of the supernatant and observed for viral CPE for 72 h. Any well demonstrating CPE within this time period was subjected to a plaque assay to determine the titer of recovered virus.

RESULTS

Preliminary complementation studies. To determine if the influenza virus HA and NA glycoproteins could substitute for the VSV G protein to generate infectious, pseudotyped particles, we initially tried to complement VSV Δ G-EGFP (30) with the HA and NA proteins of influenza virus strain A/Puerto Rico/8/34 expressed from the pCAGGS vector (49). We were able to detect only low and variable levels (10^2 to 10^3 infectious units [IU]/ml) of pseudotyped particles after complementation. Furthermore, this low level of infectivity was detectable only after trypsin treatment of the pseudotypes, indicating poor cleavage of the expressed HA protein by cellular enzymes.

Because the HA proteins of HPAIV strains are cleaved efficiently at a polybasic sequence by cellular furin-like proteases (14–17), we next tested the complementation of VSV Δ G-EGFP by an HPAIV H5 HA protein (A/Vietnam/1203/04) in combination with the PR8 (H1) NA protein. Complementation with both proteins yielded pseudotyped virus with titers of $>5 \times 10^5$ IU/ml in the absence of trypsin treatment. These pseudotype titers were only 5- to 10-fold lower than those obtained by complementation with the VSV G protein. Complementation with H5 HA or H1 NA alone did not yield infectious pseudotypes. Although HA alone could in theory confer infectivity, the NA protein is presumably required for the release of pseudotyped particles from cells through cleavage of the sialic acid receptor (50).

Construction and recovery of VSV Δ G-H5N1. Because of the positive complementation results with the H5 and N1 proteins, we next determined if a VSV Δ G recombinant expressing both the H5 HA (A/Vietnam/1203/04) and N1 (A/Puerto Rico/8/1934) proteins could be constructed and might be infectious in the absence of VSV G. We included the NA and HA genes as two separate genes with their own transcription start-stop and poly(A) signals. These genes replaced the VSV G gene, as diagrammed in Fig. 1A.

The VSV Δ G-H5N1 recombinant was recovered initially by complementation with the VSV G protein, but we found subsequently that it propagated on BHK-21 cells in the absence of the VSV G protein. The titers of VSV Δ G-H5N1 particles obtained on BHK-21 cells were typically $\sim 2 \times 10^7$ PFU per ml, or 25-fold lower than those of VSV. Consistent with the lower titers, we found that the recombinant made much smaller plaques than did VSV (Fig. 1B). A one-step growth curve of VSV Δ G-H5N1 is shown in Fig. 1C, where a maximal titer was reached by 21 h postinfection.

The VSV Δ G-H5N1 recombinant expresses both HA and NA on the cell surface. We next used indirect immunofluorescence microscopy to determine if the HA and NA proteins expressed by VSV Δ G-H5N1 were present on the cell surface. BHK-21 cells infected with VSV, VSV-H5 (a previously described VSV recombinant expressing HA from A/Hong Kong/156/97), or VSV Δ G-

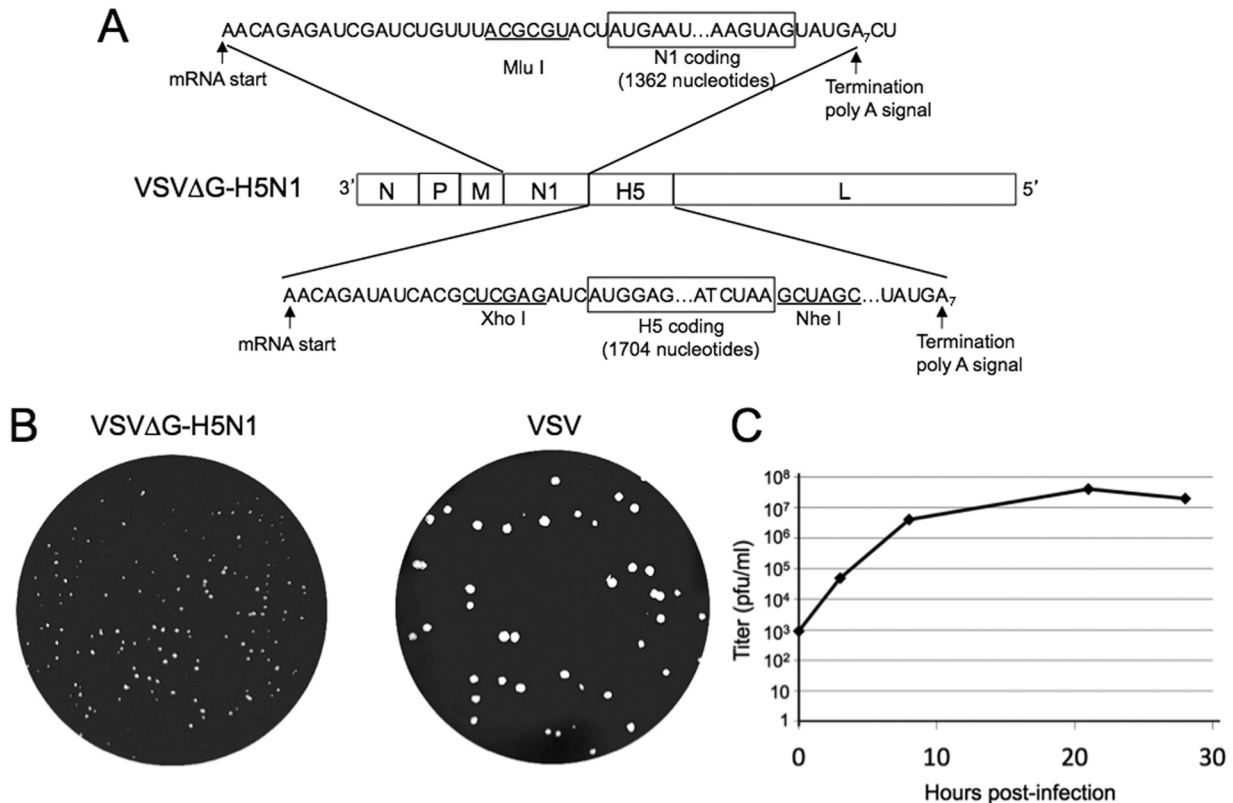


FIG 1 The VSVΔG recombinant encoding the influenza virus HA and NA genes and representative plaque morphology. (A) Diagram of the VSV recombinant with the insertion of the influenza virus N1 NA and H5 HA genes between the VSV M and L genes in a VSVΔG vector. (B) Plaque morphology of VSVΔG-H5N1 and VSV 2 days after infection of BHK-21 cells. (C) One-step growth curve of VSVΔG-H5N1. BHK-21 cells were infected at time zero with VSVΔG-H5N1 at an MOI of 5, the inoculum was removed after 30 min, cells were washed twice with PBS, and samples of the medium were taken immediately (time zero) and at the indicated times thereafter for subsequent plaque titration.

H5N1 were fixed and incubated separately with antibodies binding to VSV G, influenza virus H5 HA, or influenza virus H1 NA and then with a secondary fluorescent antibody.

The results (Fig. 2) showed that cells infected with VSVΔG-H5N1 did not express VSV G but did express H5 HA and NA on the cell surface. Control cells infected with VSV were positive for VSV G on the cell surface but negative for HA or NA. Additional control cells infected with VSV-H5 showed expression of HA and VSV G, but not NA, on the cell surface.

VSVΔG-H5N1 expresses similar levels of HA and NA. We next examined the expression levels of both HA and NA from the recombinant by metabolic labeling of infected cells with [³⁵S]methionine (Fig. 3A). BHK-21 cells were infected with VSV, VSV-H5, or VSVΔG-H5N1 and labeled for 30 min; the labeled cell lysates were then fractionated by SDS-PAGE, and the proteins were detected on a phosphorimager. The cells infected with VSV showed the major bands of all five VSV proteins (L, G, N, P, and M), while cells infected with VSVΔG-H5N1 lacked the G band and showed new protein bands with the expected mobilities of the uncleaved HA0 precursor and NA proteins. The band intensities of HA0 and NA appeared equivalent, but when corrected for methionine content (13 in HA0 and 9 in NA), the NA expression level was slightly higher, consistent with its expression from the more upstream position in the VSV genome (51). Cells infected with VSV-H5 (52) showed the HA0 precursor as well as VSV G. Note that this VSV-H5 expresses the HK156 HA protein, which mi-

grates slightly faster than the VN1203 HA protein expressed by VSVΔG-H5N1 (52).

The majority of HA incorporated into VSVΔG-H5N1 particles is cleaved to HA1 and HA2. We also used Western blotting to examine the incorporation and extent of cleavage of H5 HA present in VSV particles (Fig. 3B). Purified virus particles prepared from cells infected with VSV, VSV-H5, or VSVΔG-H5N1 were fractionated by SDS-PAGE and then blotted with anti-HA antibody or antibody to VSV. Blotting with anti-VSV antibody showed the expected major VSV proteins (G, N, and M) in VSV particles as well as the absence of VSV G in the VSVΔG-H5N1 particles. Blotting of VSVΔG-H5N1 with anti-H5 HA revealed major bands of proteins with the mobilities expected for HA1 (50 kDa) and HA2 (27 kDa) and a minor band of HA0 (77 kDa). This result indicates that the majority of HA present in the particles is cleaved. The VSV-H5 blot also showed a clear band of HA1 and a faint band of HA2, but the extent of cleavage of this HK156 H5 protein appears to be lower than that of the VN1203 protein in VSVΔG-H5N1 particles.

VSVΔG-H5N1 is nonpathogenic in mice. VSV and prototype VSV recombinants expressing VSV G and foreign antigens can cause significant pathogenesis in mice when given by the i.n. route. This pathogenesis is most easily quantified by monitoring weight loss after inoculation. To determine if VSVΔG-H5N1 was pathogenic, we inoculated BALB/c mice (5 mice per group) with 1.25×10^7 PFU of VSVΔG-H5N1 i.n. and monitored their

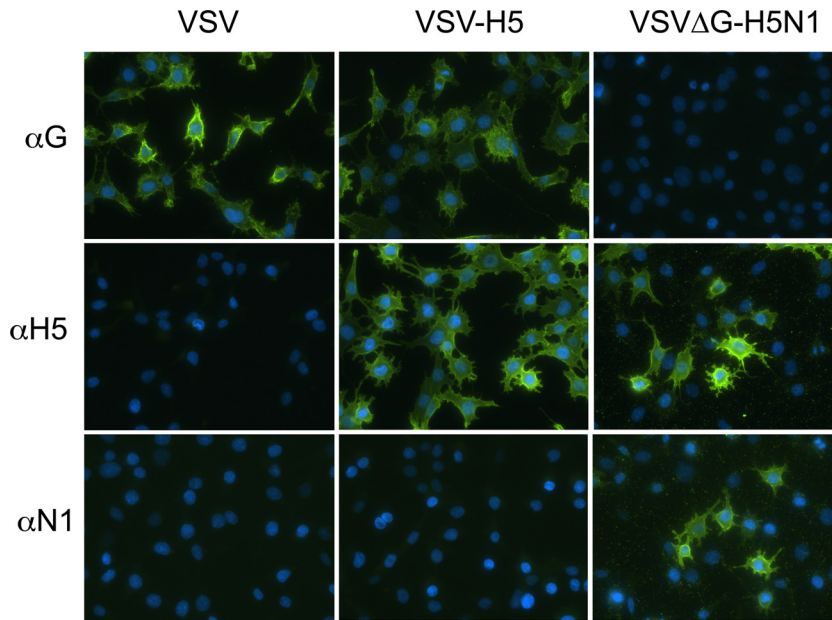


FIG 2 Indirect immunofluorescence microscopy of VSV Δ G-H5N1- and VSV-H5-infected cells. BHK-21 cells were infected with the indicated viruses for 6 h and then fixed in 3% paraformaldehyde. Cells were then incubated with either anti-VSV G mouse monoclonal antibodies, anti-H5 polyclonal sheep serum, or anti-PR8 NA polyclonal mouse serum, followed by anti-mouse or anti-sheep Alexa Fluor 488-conjugated secondary antibody staining. Nuclei are stained with DAPI. Images were taken with a Nikon Eclipse 80i fluorescence microscope.

weights for 13 days (Fig. 4). Control animals were inoculated i.n. with 10^6 PFU of a VSV recombinant expressing enhanced green fluorescent protein (VSV-EGFP) (30). The animals inoculated with VSV-EGFP showed the typical 10 to 15% weight loss at days 2, 3, and 4, followed by a gradual recovery. In contrast, the animals inoculated with VSV Δ G-H5N1 showed no weight loss. In subsequent vaccination studies, no weight loss or any other signs of pathogenesis were observed for mice inoculated with VSV Δ G-H5N1 for several months.

To determine if we could detect any replication of VSV Δ G-H5N1 *in vivo*, we inoculated mice either i.n. or i.m. with 2×10^7 PFU of VSV Δ G-H5N1 or 2×10^5 PFU of a control rVSV expressing VSV G and a hybrid influenza virus hemagglutinin (VSV-cH5/1). Tissues (leg muscle, lung, spleen, and draining lymph nodes) were isolated and homogenized at either 24 h or 96 h, as described previously, and assayed for infectious virus on BHK-21 cells (48). We detected no infectious VSV Δ G-H5N1 in any tissue at either time point. Control experiments indicated a

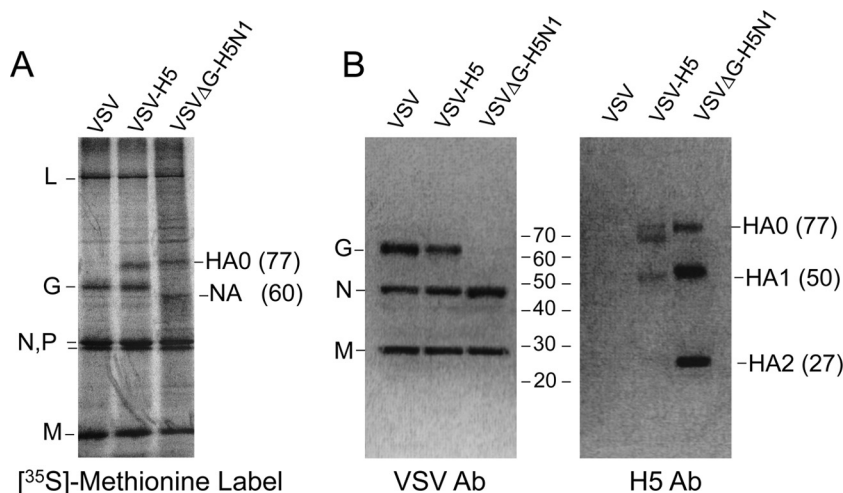


FIG 3 Metabolic labeling and Western blotting. (A) Metabolic labeling of proteins expressed from recombinant VSV vectors. BHK-21 cells were infected with the indicated viruses and then metabolically labeled with [35 S]methionine. Cell lysates were subjected to SDS-PAGE on a 4-to-12% gradient gel, and the protein bands on the dried gel were imaged. Positions of VSV proteins are indicated on the left side of the gel image, and influenza virus proteins are indicated on the right side. (B) Western blot analysis of BHK-21 whole-cell extracts infected with the indicated viruses. Sample lysates were analyzed by SDS-PAGE on a 4-to-12% gradient gel, and Western blotting was performed by using a polyclonal VSV (Indiana serotype) antiserum and the polyclonal H5 HA-specific antibody (Ab) NR-665. The positions of full-length (HA0) and cleaved (HA1 and HA2) H5 HA isoforms are indicated on the right side of the anti-H5 blot.

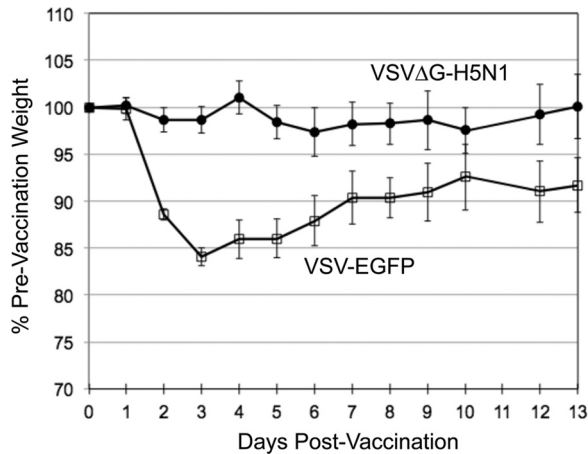


FIG 4 VSVΔG-H5N1 is nonpathogenic in mice. Groups of five 6- to 8-week-old female BALB/c mice were lightly anesthetized prior to i.n. vaccination with either 1.25×10^7 PFU of VSVΔG-H5N1 or 10^6 PFU of a VSV recombinant expressing EGFP. Weights were recorded daily for 13 days postinoculation and are graphically displayed as the average percentage of the prevaccination weight. Error bars represent 1 standard deviation above and below the average weight.

detection level of <200 PFU/tissue. In contrast, rVSV was recovered from all tissues of i.n. vaccinated mice at 24 h and from lung only at 96 h. In mice vaccinated i.m. with rVSV, virus was recovered only from muscle and draining lymph nodes. This

finding is consistent with results from previous studies using rVSVs (48). These experiments indicate that VSVΔG-H5N1 does not replicate extensively *in vivo*.

VSVΔG-H5N1 induces neutralizing antibodies to HPAIV after a single dose. To determine if VSVΔG-H5N1 could induce NAbs to HPAIV, we inoculated BALB/c mice (5 per group) with 10^6 PFU VSVΔG-H5N1 by either the i.n. or i.m. route and assayed antibody titers after 60 days using a microneutralization assay (46) for influenza virus A/Vietnam/1203/2004 carrying the homologous H5 HA. In the i.m. inoculated group, the geometric mean titer (GMT) was 39, and the titers for the individual animals varied from 28 to 101 (Fig. 5A). In the i.n. inoculated group, the GMT was 21.3, and the range was 10 to 34 (Fig. 5B). To determine if the animals developed cross-clade NAbs to influenza virus A/Indonesia/05/2005, we used pooled serum from each group because of limited material. Pooled serum from the i.m. inoculated group had a clear cross-clade Nab titer of 25, while the i.n. inoculated group had a lower titer of 10.

We also tested the neutralization of VSVΔG-H5N1 by the pooled sera obtained from the same animals at 30 and 60 days postinoculation. Serial dilutions of the sera were incubated with 100 PFU of VSVΔG-H5N1 prior to infection of BHK-21 cells. The serum dilution that gave 100% inhibition of cytopathic effect in duplicate samples was determined. As shown in Fig. 5C, the 100% neutralization titers from the i.m. inoculated group were 1:160 and 1:640 at days 30 and 60, respectively. The titers from the i.n. inoculated group were 1:80 and 1:320 at days 30 and 60, respec-

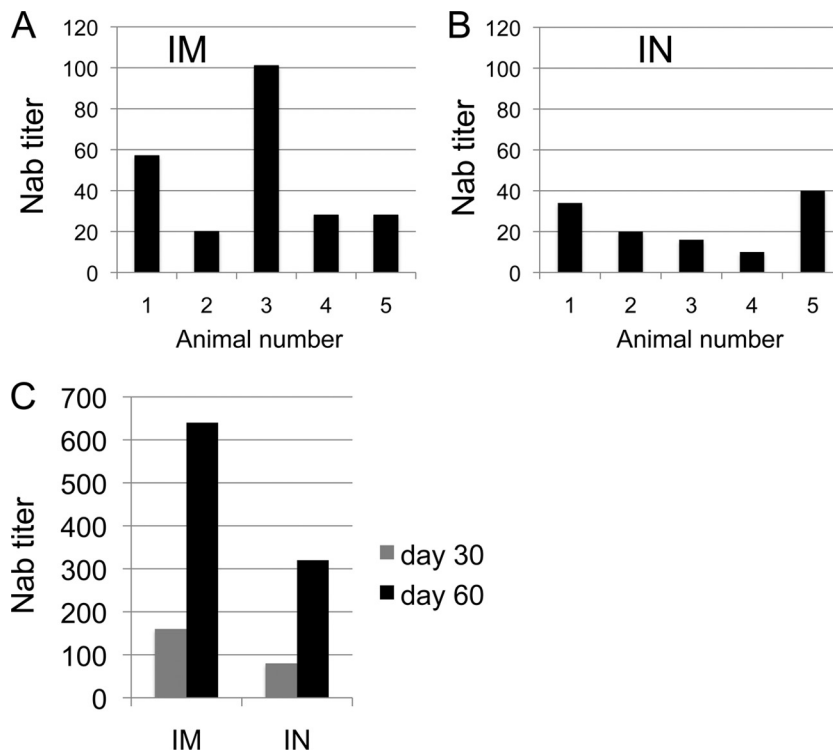


FIG 5 VSVΔG-H5N1 immunization induces NAbs in mice. Groups of five 6- to 8-week-old female BALB/c mice were vaccinated with 10^6 PFU of VSVΔG-H5N1 by either the i.m. (IM) or i.n. (IN) route, and serum was collected after 30 and 60 days to assay neutralizing antibody titers. (A) NAb titers from serum of individual mice vaccinated by the i.m. route. A microneutralization assay was performed by utilizing influenza virus A/Vietnam/1203/2004 with homologous H5 HA. (B) NAb titers from serum of individual mice vaccinated by the i.n. route as determined by a microneutralization assay. (C) Titers of NAbs in pooled 30- and 60-day-postvaccination serum from groups of 5 mice vaccinated with VSVΔG-H5N1 by either the i.m. or i.n. route. Titers are the reciprocal dilutions of pooled serum required to provide 100% neutralization of VSVΔG-H5N1 prior to incubation with BHK-21 cells.

TABLE 1 Serum from mice vaccinated with VSVΔG-H5N1 recognizes H5 HA and N1 NA but not cH6/1 HA

Mouse group	Reciprocal endpoint titer determined by ELISA								
	H5 VN04			cH6/1 (PR8 stalk)			N1 PR8		
	Expt 1	Expt 2	Avg	Expt 1	Expt 2	Avg	Expt 1	Expt 2	Avg
i.m. VSVΔG-H5N1	24,300	24,300	24,300	100	100	100	8,100	8,100	8,100
i.m. VSVΔG-G(NiV)	100	100	100	100	100	100	100	100	100

tively. The lower titers in the i.n. inoculated group were consistent with the results from the direct HPAIV NAb assays.

To better characterize the humoral response to VSVΔG-H5N1, serum collected from mice at 30 days post-i.m. vaccination with VSVΔG-H5N1 was subjected to ELISA with substrates including an H5 HA (A/Vietnam/1203/04), a chimeric HA containing an H6 HA head and an H1 HA stalk (cH6/1 HA), and N1 NA (Table 1). Endpoint titers against H5 HA and N1 NA were 1:24,300 and 1:8,100, respectively, but there was no detectable response against cH6/1 HA, indicating that the antibodies generated in response to this vaccine are largely specific for the HA head and the NA glycoprotein. The lack of reactivity with the cH6/1 HA substrate indicates a lack of heterosubtypic HA-specific and anti-stalk antibodies. Serum from mice vaccinated with VSVΔG-G(NiV) was used as a control and showed only low-level reactivity against all substrates.

Hemagglutination inhibition (HAI) experiments were performed, and mice vaccinated with a single i.m. VSVΔG-H5N1 dose did not produce detectable HAI titers 30 days after vaccination. This is not unexpected, as protective/neutralizing humoral responses against H5 HA in the absence of detectable HAI, even in human subjects, have been documented (53).

A single dose of VSVΔG-H5N1 protects mice from lethal avian influenza virus challenge. To determine if VSVΔG-H5N1 could confer protection from lethal avian influenza virus challenge, we used the same mice for which we had measured the NAb titers and challenged them i.n. with 4 50% lethal doses (LD₅₀) of the low-pathogenicity H5N1 HALo virus at 140 days after vaccination. As shown in Fig. 6A, three unvaccinated age-matched control animals showed rapid weight loss, and two of the three animals died on days 6 and 8 postchallenge. One animal made a gradual recovery to normal weight by day 22. In contrast, the five vaccinated animals in each group were all protected. On average, animals in the i.m. inoculated group lost no more than 5% of their weight, and animals in the i.n. inoculated group lost no more than 8% of their weight. Neither group showed signs of disease.

We next repeated this challenge study using a single high-dose i.m. vaccination of six mice with 1.25×10^7 PFU VSVΔG-H5N1 and a control group of six animals. The animals were challenged at 34 days postvaccination with 10 LD₅₀ of HALo. All control animals died by day 6, while all vaccinated animals were completely protected and showed no weight loss. In a separate study designed to determine how rapidly protection could be achieved, mice were vaccinated i.n. with high-dose VSVΔG-H5N1 (1.25×10^7 PFU) and challenged at 14 days postvaccination with 10 LD₅₀ of HALo. All vaccinated mice were protected from challenge (Fig. 6C), although there was a modest amount of weight loss recorded, with an average maximal weight loss of ~11% on postchallenge day 5. In contrast, control mice rapidly lost weight, and all control mice died by day 7 postchallenge.

DISCUSSION

Although there are multiple reports of pseudotyped viruses that are capable of a single round of infection after complementation with influenza virus HA (54–58), the pseudotypes are not capable of multiple cycles of replication. The recombinant VSVΔG expressing an H5 HA and an N1 NA that we describe here is replication competent in tissue culture and generates single-dose protective immunity to a lethal H5N1 challenge in mice. The use of a replication-competent vector expressing the influenza virus HA and NA proteins in place of VSV G has the major advantage of not inducing vector-specific NAb and could facilitate the use of the same vector for multiple influenza vaccines. Other VSV vector-based vaccination schemes require exchanging the G glycoprotein for that of a distinct serotype in order to avoid neutralization of boosting constructs by preexisting NAb directed at VSV G (59). VSVΔG vaccines are not subject to this limitation.

Another group recently described the construction of replication-competent VSVΔG recombinants expressing HA and NA (termed pseudotypes), but these were not tested in animals (60). As in our studies, the HA used in these hybrid VSVs was derived from a HPAIV strain containing a polybasic cleavage site. This site presumably facilitates the growth of the recombinant virus because there is efficient HA cleavage by cellular proteases.

Recombinant wild-type VSV and VSV recombinants are nonpathogenic when given by the i.m. route but can be pathogenic in mice when given by the i.n. route because they spread systemically and can cause encephalitis (33, 61, 62). Highly attenuated VSV vectors that lack detectable neurovirulence and are nonpathogenic have been developed (63–65). Initial clinical trials utilizing the attenuated VSV vector as an HIV-1 vaccine candidate in humans have shown no significant adverse events, even at very high doses (HVTN 090 study [<http://clinicaltrials.gov/ct2/show/NCT01438606>]). Furthermore, multiple VSVΔG vaccines have been shown to be nonpathogenic, even those that have *in vitro* growth capabilities restored by the addition of other viral surface glycoproteins from highly pathogenic viruses such as Ebola, Marburg, and chikungunya viruses (23, 39, 66). Given the high virulence of many HPAIV strains that have been isolated, there is concern that hybrid viruses containing components of HPAIV, especially replication-competent viruses such as VSVΔG-H5N1, may have increased virulence; however, it is clear that the expression of HA alone from HPAIVs did not confer pathogenicity to VSV in animal models, even when utilizing complete VSV-HPAIV HA strains (27, 28, 52). Our experiments additionally confirm that VSVΔG-H5N1 strains were nonpathogenic when administered i.n. to mice. We were not able to recover any VSVΔG-H5N1 from animals given this recombinant i.n. or i.m., indicating that it does not replicate to detectable

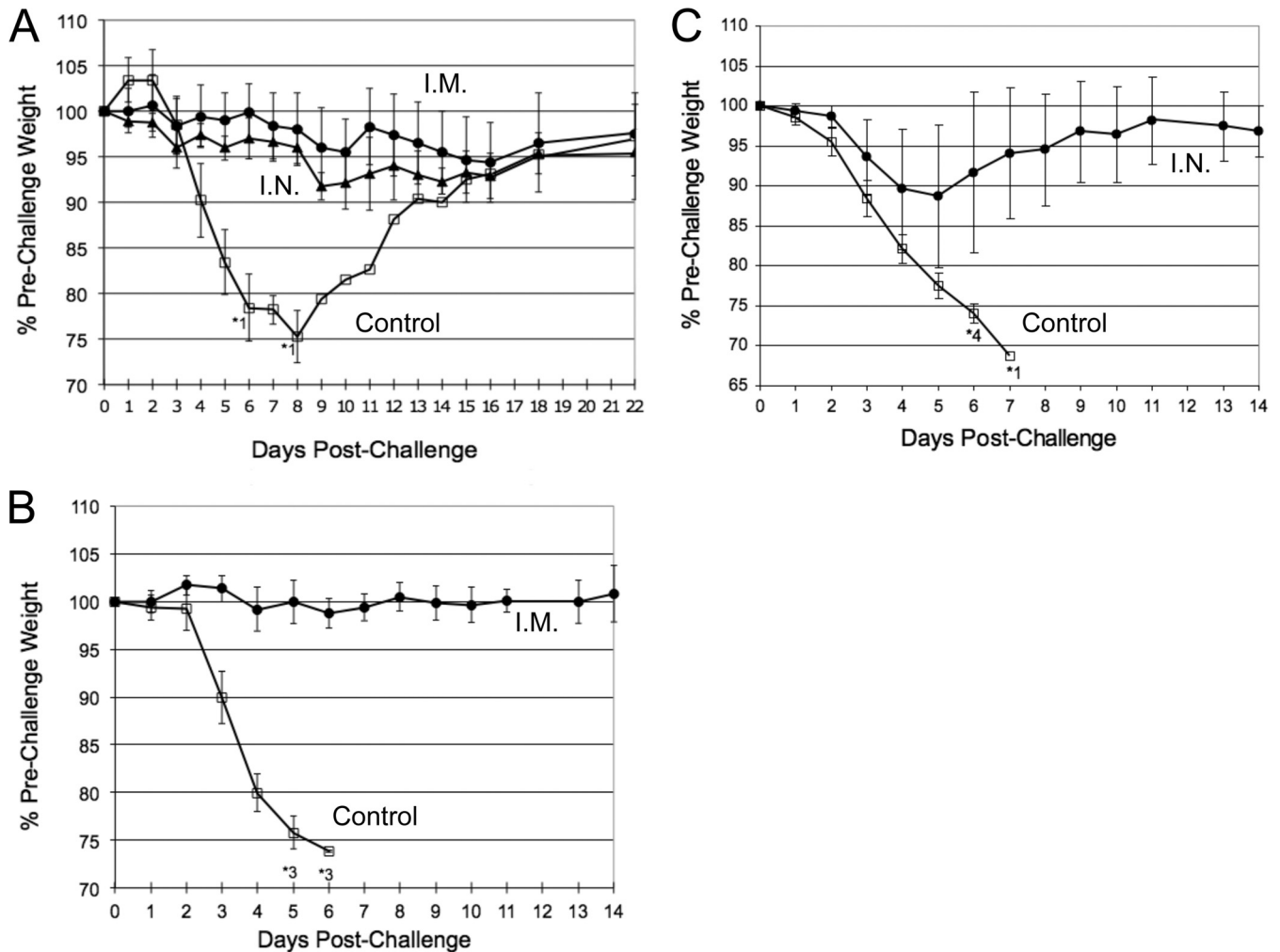


FIG 6 VSVΔG-H5N1 provides protection against H5N1 challenge. Mice vaccinated with VSVΔG-H5N1 are protected against lethal challenge with the low-pathogenicity HALo influenza challenge virus, a 6:2 gene segment reassortant virus containing HA and NA of VN1203, with the removal of the polybasic cleavage site in HA, and the internal gene segments of PR8. (A) Duration of protective immunity against HALo challenge. Groups of 5 mice vaccinated with 10^6 PFU of VSVΔG-H5N1 by either the i.m. or i.n. route were challenged with 4 LD₅₀ of HALo influenza virus at 140 days postvaccination. Control mice were a group of 3 unvaccinated age-matched mice. (B) Challenge experiments utilizing higher-dose (1.25×10^7 PFU) i.m. vaccination with VSVΔG-H5N1 and a shorter interval to challenge. Groups of 6 mice were vaccinated i.m. with 1.25×10^7 PFU of either VSVΔG-H5N1 or VSVΔG-G(NiV) (VSVΔG expressing the G glycoprotein of Nipah virus) (control group). Mice were challenged with 10 LD₅₀ of HALo influenza virus at 34 days postvaccination. (C) Short-interval challenge to establish if rapid protection against lethal challenge can be achieved. Groups of 5 mice were vaccinated i.n. with either 1.25×10^7 PFU of VSVΔG-H5N1 or 10^6 PFU of VSV-EGFP (control group). Mice were challenged with 10 LD₅₀ of HALo influenza virus at 14 days postvaccination. Error bars represent 1 standard deviation above and below the average weight. Asterisks indicate the number of animals that died at the given time points.

levels *in vivo*. Thus, this construct appears to have a novel ability to replicate efficiently *in vitro* while being nonpathogenic in an animal model.

Given the rapid spread and length of circulation of newly emergent influenza viruses, vaccine-induced protection should ideally be achieved rapidly and be of a relatively long duration. The VSVΔG-H5N1 vaccine that we describe appears to possess both of these properties: i.n. vaccination protected mice against lethal challenge after 14 and 140 days postvaccination. Additionally, the i.m. route of vaccination, which has been shown to be even more effective than i.n. vaccinations at generating both immune responses and protection in other VSVΔG models (37, 67), completely protected mice against lethal influenza virus challenge at an intermediate time point. While we anticipate that H5 HA-specific antibodies mediate protection against the

challenge virus, it is possible that there is an additional contribution from NA-specific antibodies, which were also quite robust in our experiments. Overall, these results suggest that this novel vaccine candidate is safe and effective after a single i.n. or i.m. dose. Because VSVΔG-H5N1 can be grown rapidly in tissue culture without the need for complementation, it could be produced rapidly in large quantities in response to an influenza pandemic.

Beyond its use as a vaccine candidate, the unique growth properties and apparent lack of pathogenicity of VSVΔG-H5N1 open the door to other potentially useful applications, including finding utility as a clinical and laboratory reagent for assays that would normally be precluded by biosafety level requirements. New applications for this construct as well as additional vaccination strategies are currently being explored.

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