

# Cross-Reactive Neuraminidase-Inhibiting Antibodies Elicited by Immunization with Recombinant Neuraminidase Proteins of H5N1 and Pandemic H1N1 Influenza A Viruses

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# ABSTRACT

Neuraminidase (NA), an influenza virus envelope glycoprotein, removes sialic acid from receptors for virus release from infected cells. For this study, we used a baculovirus-insect cell expression system to construct and purify recombinant NA (rNA) proteins of H5N1 (A/Vietnam/1203/2004) and pandemic H1N1 (pH1N1) (A/Texas/05/2009) influenza viruses. BALB/c mice immunized with these proteins had high titers of NA-specific IgG and NA-inhibiting (NI) antibodies against H5N1, pH1N1, H3N2, and H7N9 viruses. H5N1 rNA immunization resulted in higher quantities of NA-specific antibody-secreting B cells against H5N1 and heterologous pH1N1 viruses in the spleen. H5N1 rNA and pH1N1 rNA immunizations both provided complete protection against homologous virus challenges, with H5N1 rNA immunization providing better protection against pH1N1 virus challenges. Cross-reactive NI antibodies were further dissected via pH1N1 rNA protein immunizations with I149V (NA with a change of Ile to Val at position 149), N344Y, and I365T/S366N NA mutations. The I365T/S366N mutation of pH1N1 rNA enhanced cross-reactive NI antibodies against H5N1, H3N2, and H7N9 viruses. It is our hope that these findings provide useful information for the development of an NA-based universal influenza vaccine.

### IMPORTANCE

Neuraminidase (NA) is an influenza virus enzymatic protein that cleaves sialic acid linkages on infected cell surfaces, thus facilitating viral release and contributing to viral transmission and mucus infection. In currently available inactivated or live, attenuated influenza vaccines based on the antigenic content of hemagglutinin proteins, vaccine efficacy can be contributed partly through NA-elicited immune responses. We investigated the NA immunity of different recombinant NA (rNA) proteins associated with pH1N1 and H5N1 viruses. Our results indicate that H5N1 rNA immunization induced more potent cross-protective immunity than pH1N1 rNA immunization, and three mutated residues, I149V, I365T, and S366N, near the NA enzyme active site(s) are linked to enhanced cross-reactive NA-inhibiting antibodies against heterologous and heterosubtypic influenza A viruses. These findings provide useful information for the development of an NA-based universal influenza vaccine.

Members of the Orthomyxoviridae family, influenza A viruses are enveloped viruses containing a single-strand, 8-segment, negative-sense RNA genome typically encoding 11 or 12 viral proteins (1, 2). Influenza A virus subtypes have been classified based on the antigenic properties of hemagglutinin (HA) and neuraminidase (NA) glycoproteins, designated H1 to H16 and N1 to N9, respectively. Besides these subtypes, H17N10 and H18N11 viruses from fruit bats were also identified (3, 4). According to phylogenetic analyses, subtypes N1 to N9 can be classified as belonging to group 1 (including N1, N4, N5, and N8) or group 2 (including N2, N3, N6, N7, and N9) (5). To date, N1, N2, N7, and N9 subtypes are known to trigger human epidemics (6, 7). N3 and N8 subtypes only cause limited sporadic human infections, as reported for H7N3 in Mexico (8) and Canada (9) and for H10N8 in China (10).

NA, an enzymatic protein with a complex tetrameric structure (11), is capable of cleaving sialic acid linkages on cell surfaces, thereby facilitating viral release from infected cells (12). NA also contributes to viral transmission and infection by destroying decoy receptors on cilia, mucins, and cellular glycocalyx (12–15). NA immunogenicity was first observed in human subjects immunized with an NA-specific inactivated vaccine (16). The use of recombinant NA (rNA) proteins expressed in yeast (17) or insect cells (18) elicits protection against lethal virus challenges in im-

munized mice. Ferrets immunized with rNA proteins exhibit a distinctive type of protection in addition to that provided by HA immunization alone (19). NA-inhibiting (NI) antibodies are known to limit virus spreading and to mitigate clinical symptoms of influenza A virus infection (19). Mice immunized with a reverse-genetic reassortant H1N1 virus containing seasonal influenza virus NA exhibit cross-reactive NI antibodies and reduced mortality from pandemic H1N1 (pH1N1) virus challenges (20). Live attenuated influenza vaccines for seasonal H1N1, H3N2, and pH1N1 strains have been reported as inducing cross-reactive NI antibodies to H5N1 viruses in ferrets (21), and NI antibodies elic-

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FIG 1 Expression and characterization of soluble recombinant NA (rNA) proteins. The cDNA coding sequences of NA ectodomains (A/Vietnam/1203/2004 [H5N1] or A/Texas/05/2009 [pH1N1]) with additional N-terminal sequences of 6×His residues and hvsp sequences were constructed and expressed using a baculovirus-insect cell expression system. (A and B) Purified rNA proteins (2 μg) were confirmed by Coomassie blue staining (A) and Western blotting (B). (C) Fetuin-based assays were used to determine rNA protein enzymatic activity.

ited by a seasonal trivalent influenza vaccine have been reported as providing cross-protective immunity against lethal H5N1 challenges, also in ferrets (22). Furthermore, NA-based virus-like particles (VLPs) containing NA and matrix proteins M1 and M2 have been shown to elicit more potent NI antibodies and to confer cross-protective immunity against H5N1 and pH1N1 viral challenges in mice (23). NI antibodies have also been detected in humans vaccinated with an H5N1 inactivated vaccine (24), as well as in humans exposed to natural infections (25). NA immunogenicity and cross-protection mechanisms remain unclear.

For this study, we used a baculovirus-insect cell expression system to obtain soluble rNA proteins from H5N1 and pH1N1. Two-dose active immunizations in mice were used to evaluate the immunogenicities of H5N1 rNA and pH1N1 rNA. In addition to detecting NA-specific total IgG and IgG subtypes, NI antibodies in sera, and antibody-secreting B cells (ASCs) in splenocytes, we also observed protective immunity following live-virus challenges. Our data indicate that H5N1 rNA and pH1N1 rNA immunizations provided complete protection against homologous virus challenges, with H5N1 rNA immunization eliciting more potent cross-reactive NI antibodies and inducing better protection against heterologous pH1N1 virus challenges. The residues at positions 365 and 366 near the NA enzyme active site were observed associating with enhanced cross-reactive NI antibodies against heterologous and heterosubtypic influenza viruses. It is our hope that these findings provide useful information for the development of an NA-based universal influenza vaccine.

# MATERIALS AND METHODS

**Recombinant NA protein expression and purification.** cDNA from the NA genes of A/Vietnam/1203/2004 (H5N1) (GI 145284408) and A/Texas/05/2009 (pH1N1) (GI 255602223) was separately synthesized with insect cell-optimized codon sequences from Genomics, Inc. The coding sequences of the NA H5N1 and pH1N1 ectodomains, together with additional N-terminal sequences containing gp67 signal peptides (MLLVNQ SHQGFNKEHTSKMVSAIVLYVLLAAAAHSAFA), six-His residues (HH HHHH), tetrameric human vasodilator-stimulated phosphoprotein

(hvsp) domains (SSSDYSDLQRVKQELLEEVKKELQKVKEEIIEAFVQE LRKRGS), and a thrombin cleavage site (LVPRGS), were cloned into pFastBac expression vectors (26). Next, rNA proteins were produced using a Bac-to-Bac insect cell expression system (Invitrogen) according to the manufacturer's instructions. Briefly, Sf9 cells were infected with recombinant baculoviruses expressing the NA ectodomains of H5N1 and pH1N1 for 48 h prior to collecting supernatants for additional rNA protein purification using nickel-chelated resin affinity chromatography (Tosoh). H5N1 rNA and pH1N1 rNA purity was confirmed by Coomassie blue staining. Anti-His horseradish peroxidase (HRP)-conjugated antibodies (Affymetrix) were used for Western blotting characterization.

Production and purification of H5N1 and H7N9 VLPs. H5N1 and H7N9 VLPs were produced using the procedures described in reference 27. Briefly, the H5 HA gene of A/Thailand/1(KAN-1)/2004 (H5N1), the H7 HA gene of A/Shanghai/2/2013 (H7N9), and the M1 gene of A/WSN/ 1933 (H1N1) were cloned into separate pFastBacDual vectors (Invitrogen). The N1 NA gene of A/Vietnam/1203/2004 (H5N1), the N9 NA gene of A/Shanghai/2/2013 (H7N9), and the M2 gene of A/WSN/1933 (H1N1) were cloned into separate vectors. For H5N1 VLP production, Sf9 cells were coinfected with Bac-H5HA-M1 and Bac-N1NA-M2 recombinant baculovirus at a multiplicity of infection (MOI) of 3 or 1, respectively. Culture supernatants were harvested and concentrated 72 h postinfection. VLPs were further purified using a 20% sucrose solution and centrifugation at 33,000 rpm for 3 h. H5N1 VLPs were obtained and stored at 4°C until used for NI assays. For H7N9 VLPs, Sf9 cells were coinfected with Bac-H7HA-M1 and Bac-N9NA-M2 recombinant baculovirus at an MOI of 3 or 1, respectively, for 72 h. The subsequent steps were the same as for H5N1 VLP production.

Mouse immunizations. BALB/c mice (6 to 8 weeks old) purchased from the Taiwan National Laboratory Animal Center were immunized twice intramuscularly with a 3-week interval with 2 or 20  $\mu$ g of rNA proteins plus 10  $\mu$ g CpG and 10% poly(ethylene glycol)-block-poly(lactide-co-epsilon-caprolactone) (PEGb-PLACL), squalene, and Span85 (PELC) emulsion as described in a previous report (28). Serum samples were collected 2 weeks after the second inoculation; splenocytes were harvested and isolated 1 week later. Procedures involving live animals were performed according to guidelines established by the Laboratory Animal Center of National Tsing Hua University (NTHU). All animal use



FIG 2 NA-specific IgG antibodies (Ab) against different viruses induced by rNA immunization. Female BALB/c mice were intramuscularly immunized with 2 or 20  $\mu$ g of H5N1 rNA or pH1N1 rNA plus 10% PELC and 10  $\mu$ g CpG adjuvant twice with a 3-week interval; serum samples were collected 14 days after the second inoculation. Briefly, 96-well plates were coated with 2  $\mu$ g/ml of recombinant N1 NA proteins (A), 10  $\mu$ g/ml of H5N1 VLPs (B), 20  $\mu$ g/ml of the pH1N1 virus (C), 20  $\mu$ g/ml of the H3N2 virus (D), or 10  $\mu$ g/ml of H7N9 VLPs (E) and held overnight at 4°C. ELISAs with H5N1 rNA or pH1N1 rNA mouse antisera or anti-mouse N1 NA-specific antibodies (Abcam) were performed to measure NA-specific total IgG titers (0.2 endpoint). Data are expressed as means  $\pm$  standard deviations (SD) for three experiments.

protocols were reviewed and approved by the NTHU Institutional Animal Care and Use Committee (IACUC) (approval no. 10002).

**Viral challenges.** For the two-dose immunization strategy, BALB/c mice (6 to 8 weeks old) were placed in one of 5 groups, with each group consisting of 5 mice immunized twice with a 3-week interval with either 2 or 20  $\mu$ g of H5N1 rNA or pH1N1 rNA proteins plus CpG and PELC or with phosphate-buffered saline (PBS). Three weeks after the second inoculations, all mice were intranasally challenged with 10 50% murine lethal doses (MLD<sub>50</sub>) of the H5N1 (NIBRG-14 [RG-14]), pH1N1 (A/California/ 07/2009 [CA/09]), or H7N9 (A/Taiwan/01/2013 [TW/13]) virus. PBS-immunized mice were used as a mock control. Survival rates and body weights were recorded daily for 14 days. For mouse virus challenges, the experimental procedures were reviewed and approved by the IACUC of Academia Sinica, Taiwan. According to IACUC guidelines, a weight loss of 25% or more was established as an endpoint.

**Enzyme-linked immunosorbent assays (ELISAs).** Individual wells in 96-well plates were coated with purified rNA proteins (100  $\mu$ l at 2  $\mu$ g/ml) and held overnight at 4°C, washed 3 times with PBST (0.05% Tween 20 in PBS), and blocked with blocking buffer (1% bovine serum albumin [BSA] in PBS) for at least 1 h. Next, 100  $\mu$ l of 2-fold serially diluted serum samples were added and held at room temperature (RT) for 1 h, followed

by 3 additional washes with PBST. HRP-conjugated goat anti-mouse IgG antibodies (Bethyl Laboratories, Inc.) were added to each well, the plates were incubated for 1 h, and the wells washed 3 times with PBST. Anti-NA IgG titers were determined by adding TMB substrate (3,3',5,5'-tetra-methyl benzidine; Biolegend), holding for 15 min at RT, and stopping the reaction with 2 N  $H_2SO_4$ . Endpoint titers were determined as the reciprocals of the most-diluted serum concentrations giving a mean optical density (OD) at 450 nm of more than 0.2.

NI assay. To determine the optimal concentrations of viruses or VLPs for the subsequent NI assays, the NA activities induced by each virus or VLPs as determined from ELISA data were plotted in GraphPad Prism 5.0 and fit to a nonlinear curve as previously described (29). The amounts of each virus or VLP reacted with fetuin and agglutinin resulted in OD values of around 2.0, half the maximal OD value in the ELISAs. Quantities of 1  $\mu$ g of VLPs (H5N1 or H7N9) or 10<sup>5</sup> PFU of viruses (pH1N1 or H3N2) were used for NI serological assays. NI antibodies were measured using a previously described fetuin-based assay procedure (30). Briefly, 96-well plates were coated with 50  $\mu$ g/ml fetuin (Sigma) and held overnight at 4°C before being washed 3 times with PBST and blocked with blocking buffer for 2 h. Twofold serially diluted serum samples in blocking buffer were incubated with equal volumes containing 1  $\mu$ g VLPs (H5N1 or H7N9) or



FIG 3 NA-inhibiting-antibody curves and the corresponding IC<sub>50</sub>s against different virus strains produced by immunizing mice with H5N1 rNA and pH1N1 rNA. Fetuin-based assays were used to measure NA-inhibiting-antibody titers as reductions in the NA enzymatic activities of the pH1N1 or H3N2 viruses or H5N1 or H7N9 VLPs. (A to H) H5N1 rNA inhibition curves against homologous H5N1 VLPs (A), heterosubtypic pH1N1 viruses (B), the H3N2 viral strain (C), or H7N9 VLPs (D) are plotted. pH1N1 rNA inhibition curves are plotted against the homologous pH1N1 viral strain (E), heterosubtypic H5N1 VLPs (F), the H3N2 viral strain (G), or H7N9 VLPs (H). (I) Corresponding IC<sub>50</sub>s were determined as the 50% reduction in the NA enzymatic activities of the homologous strains (white bars) or heterosubtypic viruses (filled bars). Integrated data are expressed as means  $\pm$  SD from three experiments. \*, P < 0.05; \*\*, P < 0.01.

10<sup>5</sup> PFU viruses (pH1N1 or H3N2) for 1 h at 37°C, added to the fetuincoated plates, held for another 1 h at 37°C, and then washed 3 times with PBST. Peroxidase-labeled peanut agglutinin (100  $\mu$ l at 2.5  $\mu$ g/ml) (Sigma) was added to each well, incubated for 1 h at RT, and washed 3 times with PBST. The NA activity levels of viruses (pH1N1 or H3N2) and VLPs (H5N1 or H7N9) were determined by adding TMB substrate (Biolegend), holding for 15 min at RT, and stopping the reaction with 2 N H<sub>2</sub>SO<sub>4</sub>. Plates were read with an ELISA reader (Tecan) at an OD of 450 nm. The corresponding 50% inhibitory concentrations (IC<sub>50</sub>s) were defined as the reciprocal serum dilutions inhibiting 50% of viral NA enzyme activity.

NA-specific antibody-secreting B cells. Splenocytes were collected from each group of rNA-immunized or PBS-immunized mice 3 weeks after the second inoculations. Multiscreen 96-well filtration plates (Millipore) were coated with rNA proteins (1  $\mu$ g per well) and incubated overnight at 4°C. Plates were blocked with 200  $\mu$ J/well of complete RPMI 1640 (10% fetal bovine serum [FBS], 1× penicillin/streptomycin, 1× sodium pyruvate, 1× nonessential amino acids [NEAA], and 100  $\mu$ M  $\beta$ -mercaptoethanol) and held for 1 h at RT. Splenocytes (2 × 10<sup>5</sup>) diluted in complete RPMI 1640 were added to individual plates and incubated for 48 h at 37°C. After 3 washes with PBST, HRP-conjugated anti-mouse IgG was added to each well and held for 2 h at RT. After 3 PBST and 2 PBS washes, 3-amino-9-ethylcarbazole (AEC) substrate was added to each plate and the plates held at RT for approximately 10 to 60 min before the reactions were stopped with double-distilled water. Immunospots for each immunized group were determined using an enzyme-linked immunosorbent spot assay (ELISPOT) plate reader (CTL, Inc.).

**Statistical analysis.** All results were analyzed using two-tailed Student's *t* tests. Asterisks in the figures indicate statistical significance as follows: \*, P < 0.05; \*\*, P < 0.01; and \*\*\*, P < 0.001.

## RESULTS

H5N1 rNA and pH1N1 rNA proteins were expressed and purified using a baculovirus-insect cell expression system. Soluble H5N1 rNA and pH1N1 rNA were produced from the culture supernatants of Sf9 insect cells infected by recombinant baculoviruses and purified using Ni-chelated affinity chromatography, and their production yields were approximately 0.5 mg/liter and 0.25 mg/liter, respectively, with 80 to 90% purity (Fig. 1A). According to the results from SDS-PAGE gels with Coomassie blue staining (Fig. 1A) and Western blotting (Fig. 1B), purified H5N1 rNA and pH1N1 rNA proteins had molecular masses of 53 kDa each. Based on Eadie-Hofstee measurements, the  $K_m$  values of H5N1 rNA and pH1N1 rNA were 2.80 and 1.90, respectively (Fig. 1C), indicating enzyme activity of insect cell-expressed H5N1 rNA and pH1N1 rNA.

NA-specific IgG antibodies were induced by H5N1 rNA and pH1N1 rNA immunizations. Groups of five female BALB/c mice were immunized intramuscularly with two doses of H5N1 rNA or pH1N1 rNA proteins (2 or 20 µg per dose) with a 3-week interval between doses. Antisera were collected 2 weeks after the second immunization. Data for NA-specific IgG titers against the same immunogens (H5N1 rNAs or pH1N1 rNA proteins) in each group are presented in Fig. 2A. We used an anti-mouse N1 NAspecific antibody as a positive control and PBS-immunized sera as a negative control (NA-specific IgG titers were 10<sup>3.5</sup> to 10<sup>4</sup> and undetectable, respectively). As shown by the results in Fig. 2A, mice immunized with 20 µg of H5N1 rNA or pH1N1 rNA exhibited slightly higher NA-specific total IgG titers than mice immunized with 2 µg of H5N1 rNA or pH1N1 rNA. We also performed ELISAs with H5N1 VLPs (Fig. 2B) and the pH1N1 (A/California/ 04/2009) virus (Fig. 2C). Our data indicate that at either 2 or 20 µg, both H5N1 rNA and pH1N1 rNA cross-reacted with H5N1 VLPs and the pH1N1 virus to induce similar total IgG titers (Fig. 2B and C). Similar cross-reactive results were also found for the H3N2 virus (A/Udorn/307/1972) and H7N9 VLPs (Fig. 2D and E). No significant differences were noted for either IgG1 or IgG2a subclass titers in the same immunization groups. The single exception was significantly greater IgG2a for mice immunized with 20 µg H5N1 rNA (data not shown).

NI antibodies are elicited by H5N1 rNA and pH1N1 rNA immunizations. To measure NI antibody titers, 2-fold serially diluted serum samples from each immunization group were mixed with 1 µg of H5N1 VLPs, 10<sup>5</sup> PFU of the pH1N1 (A/California/ 04/2009) virus, 10<sup>5</sup> PFU of the H3N2 (A/Udorn/307/1972) virus, or 1 µg of H7N9 VLPs and then examined using fetuin-based assays. The percentages of NA inhibition and NI antibody titers against H5N1 and pH1N1 in each immunization group are shown in Fig. 3A, B, E, and F. Those results indicate dose dependency, with the NI curves from all rNA immunization groups being significantly higher than the curves for the PBS immunization group. NA inhibition percentages and NI antibody titers were also demonstrated for the H3N2 virus and H7N9 VLPs, with similar results but lower NI antibody titers (Fig. 3C, D, G, and H). Regarding the IC<sub>50</sub>s, similar NI antibody titer ranges were observed against the homologous viruses, 3.7 to 3.8 for H5N1 rNA immunization against H5N1 and 4.3 to 4.4 for pH1N1 rNA immunization against pH1N1 (Fig. 3I). However, significant differences were noted among the six immunization groups in terms of NI antibody titers against the H5N1, pH1N1, H3N2, and H7N9 heterologous viruses. Both H5N1 rNA immunization groups (2 and 20 µg) expressed higher heterologous NI antibody titers against pH1N1 than were elicited by the two pH1N1 rNA immunization groups against H5N1 viruses. In contrast, H5N1 rNA and pH1N1 rNA elicited lower NI antibody titers, with IC<sub>50</sub>s of less than 2.32 for all groups immunized against the H3N2 and H7N9 viruses (Fig. 3I). Our data indicate that the effect of H5N1 rNA immunization was limited to more potent heterologous NI antibodies against pH1N1 viruses.



FIG 4 Detection of antibody-secreting B cells in mouse spleens induced by H5N1 rNA and pH1N1 rNA proteins. ELISPOT assays were used to measure IgG-secreting B cell numbers. Splenocytes collected from mice immunized with H5N1 rNA ( $2 \times 10^5$ ) were cocultured with 1 µg of H5N1 rNA or pH1N1 NA proteins and held for 48 h at 37°C. Splenocytes from mice immunized with pH1N1 rNA were incubated for 48 h at 37°C with 1 µg of pH1N1 rNA or H5N1 NA proteins. Immunospots were determined by ELISPOT assays. Data are expressed as the means ± SD from at least two experiments. \*, P < 0.05.

Antibody-secreting B cells detected in spleen. To measure anti-NA IgG-secreting B cells elicited by H5N1 rNA or pH1N1 rNA, splenocytes were collected from immunized mice 3 weeks after their second immunization, reacted with 1  $\mu$ g H5N1 rNA or pH1N1 rNA protein per well, and examined using ELISPOT assays. The results shown in Fig. 4 indicate that the numbers of spots against homologous viruses due to H5N1 rNA immunization were slightly higher than those resulting from pH1N1 rNA immunization. Significantly higher numbers of spots against H5N1 and pH1N1 heterologous viruses were only noted in the 20- $\mu$ g H5N1 rNA immunization group, which also exhibited higher quantities of ASCs in splenocytes against homologous and heterologous H5N1 and pH1N1 viruses.

Protective immunity against H5N1, pH1N1, and H7N9 virus challenges. To assess protective immunity triggered by rNA immunizations, mice immunized with 2 or 20 µg of H5N1 rNA or pH1N1 rNA proteins were challenged with 10 MLD<sub>50</sub> of H5N1 (RG-14), pH1N1 (CA/09), or H7N9 (TW/13) viruses 3 weeks after their second immunizations. According to the results shown in Fig. 5A and B, with the exceptions of an 80% survival rate for mice immunized with 2 µg of pH1N1 rNA and a 0% rate for the PBS control mice, all immunization groups had 100% survival rates after homologous H5N1 or pH1N1 viral challenges. Significantly smaller body weight losses were observed for mice in the 20-µg H5N1 rNA and 20-µg pH1N1 rNA immunization groups challenged with homologous H5N1 or pH1N1 than for those in the 2-µg H5N1 rNA and 2-µg pH1N1 rNA immunization groups (Fig. 6A and B). In terms of cross-protection levels, mice receiving either 2- or 20-µg H5N1 rNA inoculations exhibited complete protection against pH1N1 viral challenges (Fig. 5C), with significantly smaller body weight losses noted in the 20-µg group



FIG 5 H5N1 rNA and pH1N1 rNA protective immune responses against different virus challenges. (A and C) Mice were intramuscularly immunized with 2 or 20  $\mu$ g of H5N1 rNA twice with a 3-week interval and intranasally challenged with 10 MLD<sub>50</sub> of the homologous H5N1 (RG-14) virus (A) or heterologous pH1N1 (CA/09) (C) at the end of week 6. Shown are survival curves for mice immunized with H5N1 rNA over a 14-day period after challenge. (B and D) Mice immunized with 2 or 20  $\mu$ g of pH1N1 rNA were challenged with 10 MLD<sub>50</sub> of the homologous pH1N1 (CA/09) virus strain (B) or heterologous H5N1 (RG-14) virus (D) at the end of week 6. Shown are survival curves for mice immunized with pH1N1 rNA over a 14-day period after challenge. (E) Mice were intramuscularly immunized with 2 or 20  $\mu$ g of pH1N1 rNA were challenged with 10 MLD<sub>50</sub> of the homologous pH1N1 (CA/09) virus strain (B) or heterologous H5N1 (RG-14) virus (D) at the end of week 6. Shown are survival curves for mice immunized with pH1N1 rNA over a 14-day period after challenge. (E) Mice were intramuscularly immunized with 20  $\mu$ g of H5N1 rNA or pH1N1 rNA twice with a 3-week interval and intranasally challenged with 10 MLD<sub>50</sub> of heterosubtypic H7N9 (TW/13) viruses 3 weeks after the second inoculations. Survival rates over the 14 days after the viral challenges are shown.

(Fig. 6C). A 60% survival rate (Fig. 5D) and faster weight recovery following challenge with the heterologous H5N1 virus were observed in the 20- $\mu$ g pH1N1 rNA immunization group (Fig. 6D). No protection against the heterosubtypic H7N9 virus challenge was observed in the 20- $\mu$ g H5N1 rNA and 20- $\mu$ g pH1N1 rNA immunization groups (Fig. 5E and 6E).

**Cross-reactive NI antibodies elicited by pH1N1 rNA mutant proteins.** To add detail to our investigation of cross-reactive NI epitopes, we aligned the amino acid sequences of A/Vietnam/ 1203/2004 (H5N1), A/Texas/05/2009 (pH1N1), A/Udorn/307/ 1972 (H3N2), and A/Shanghai/02/2013 (H7N9), as shown in Fig. 7. Based on the alignment analysis, we identified 34 different amino acids (Fig. 7, red highlighting) in the NA ectodomains of A/Vietnam/1203/2004 (H5N1) and A/Texas/05/2009 (pH1N1) sequences (94.6% identical). Previous reports have identified influenza NA enzyme catalytic sites at residues 118 and 119, 151 and 152, 198, 224, 227, 243, 274, 276 and 277, 292, 330, 350, and 425 (11, 30); therefore, we targeted residues 149, 344, 365, and 366, all located close to NA enzyme active sites and contributing to NI

antibody elicitation. These four target residues (with changes of Ile to Val at position 149 [I149V], N344Y, and I365T/S366N) are shaded in blue and the enzyme active sites are shown as red sticks in a three-dimensional (3-D) NA structure diagram in Fig. 8. Sitedirected mutagenesis at these four residues produced the three mutant pH1N1 rNA proteins with mutations I149V, N344Y, or I365T/S366N (Fig. 9). The results from an analysis of antisera from the wild type and the three mutant pH1N1 rNAs show that the I149V and I365T/S366N proteins elicited more NI antibodies against the homologous pH1N1 strain (Fig. 10A), with the I365T/S366N protein eliciting more potent cross-reactive NI antibodies against the H5N1 (Fig. 10B), H3N2 (Fig. 10C), and H7N9 viruses (Fig. 10D). The corresponding IC<sub>50</sub>s calculated from NI response curves indicate that the I149V and I365T/S366N mutant proteins resulted in increased NI antibody titers against the homologous pH1N1 viruses and that all three mutant proteins resulted in increased NI antibody titers against heterosubtypic H3N2 and H7N9 viruses (Fig. 10E). The I365T/S366N mutation of pH1N1 rNA induced the highest NI



FIG 6 Body weight recovery tied to H5N1 rNA and pH1N1 rNA immunization responses against different virus challenges. (A and C) Mice were intramuscularly immunized with 2 or 20  $\mu$ g of H5N1 rNA twice with a 3-week interval and intranasally challenged with 10 MLD<sub>50</sub> of the homologous H5N1 (RG-14) virus (A) or heterologous pH1N1 (CA/09) (C) at the end of week 6. Shown are body weight losses over a 14-day period for mice immunized with H5N1 rNA. (B and D) Mice immunized with 2 or 20  $\mu$ g of pH1N1 rNA were challenged with 10 MLD<sub>50</sub> of the homologous pH1N1 (CA/09) virus strain (B) or heterologous H5N1 (RG-14) virus (D) at the end of week 6. Shown are body weight losses over a 14-day period for mice immunized with H5N1 rNA. (B and D) Mice immunized with 2 or 20  $\mu$ g of pH1N1 rNA were challenged with 10 MLD<sub>50</sub> of the homologous pH1N1 (CA/09) virus strain (B) or heterologous H5N1 (RG-14) virus (D) at the end of week 6. Shown are body weight losses over a 14-day period for mice immunized with pH1N1 rNA. (E) Mice were intramuscularly immunized with 20  $\mu$ g of H5N1 rNA or pH1N1 rNA twice with a 3-week interval and intranasally challenged with 10 MLD<sub>50</sub> of heterosubtypic H7N9 (TW/13) viruses 3 weeks after the second inoculation. Body weights were measured for the 14 days after each viral challenge. Body weight loss of >25% was used as an endpoint.

antibody titers against the homologous, heterologous, and heterosubtypic strains.

#### DISCUSSION

NA-based influenza vaccines are attractive because of the relatively smaller number of changes in NA antigens compared to the changes in HA antigens in host immune systems (31, 32). For this study, we constructed and purified H5N1 rNA and pH1N1 rNA proteins from Sf9 insect cells and found that mice immunized with H5N1 rNA and pH1N1 rNA proteins exhibited higher quantities of NA-specific total IgG, IgG1, IgG2a subclass, and NI antibodies, increased numbers of ASCs in the spleen, and better protective immunization induced more potent cross-reactive NI antibodies and protective immunity against pH1N1 viruses than did pH1N1 rNA immunization against H5N1 viruses. Cross-reactive NI epitopes were further dissected by immunization using pH1N1 rNA proteins with I149V, N344Y, and I365T/S366N NA mutations. The I365T/S366N mutation of pH1N1 rNA was found to increase cross-reactive NI antibodies against heterologous and heterosubtypic influenza A viruses.

H5N1 rNA and pH1N1 rNA proteins expressed in baculovirus-infected Sf9 cells and purified by affinity chromatography displayed similar enzymatic activities. The  $K_m$  value of pH1N1 rNA was slightly lower (higher affinity) than that of H5N1 rNA (Fig. 1C). Note that only NA ectodomain proteins (amino acids [aa] 62 to 447 for H5N1 and aa 82 to 467 for pH1N1) were encoded for soluble rNA expression. The coding sequences did not include N-terminal stalk regions; that is, H5N1 lacked aa 38 to 57 and pH1N1 lacked aa 38 to 77 (Fig. 7). Each NA monomer consists of six identical  $\beta$  sheets ( $\beta$ ) plus several strands (S) and loops (L). Influenza NA enzyme catalytic sites are located at the upper corners of each monomer (tetrameric structure); amino acids encircling active catalytic sites are highly conserved among various influenza A and B virus strains (11, 33). A comparison of the NA sequences of H5N1 (A/Vietnam/1203/2004) and pH1N1 (A/Texas/05/2009), based on X-ray crystallographic diffraction data for NA active sites from A/Tokyo/3/67, did not identify

	1	1 Enzyme active site(s)					
A/Texas/05/2009(H1N1)	MNPNQKIITI	GSVCMTIGMA	NLILQIGNII	SIWISHSIQL	GNQNQIETCN	QSVITYENNT	
A/Vietnam/1203/2004 (H5N1)		••I••VT•IV	S•M•••A•M•	•••V•••HT	•••H•S•P		
A/Udorn/307/1972 (H3N2)		•••SL••ATI	CFLM••AILV	TTVTLHF	KQYECDSPA•	NQ•MPC•PII	
A/Shanghai/02/2013 (H7N9)	•••••LCT	SATAII••AI	AVLIGMA•LG	LNIGLH•	KP-GCNCSHS	• PET • NTSQ •	
	61						
A/Texas/05/2009(H1N1)	WVNQTYVNIS	NTNFAAGQSV	VSVKLAGNS-	SLCP <mark>VS</mark> GW	AIYSKDNSVR	IGSKGDVFVI	
A/Vietnam/1203/2004 (H5N1)	••	••••LTEKA•	A•••••-	•• <b>IN</b> ••	•v••••1•	•••••	
A/Udorn/307/1972 (H3N2)	IERNITEIVY	L••TTIEKEI	CPKLVEYRNW	SKPQ•KIT•F	• PF••••I•	LSAG••IW•T	
A/Shanghai/02/2013 (H7N9)	II•NY•NETN	I • • IQMEERT	SRNF•NL	TKG••TINS•	H••G•••AV•	••ESS••L•T	
	121			149			
A/Texas/05/2009(H1N1)	REPFISCSPL	ECRTFFLTQG	ALLNDKHSNG	TIKDRSPYRT	LMSCPIGEVP	SPYNSRFESV	
A/Vietnam/1203/2004 (H5N1)	••••• <u>H</u> •			•v••••H••	••••V••A•	•••••	
A/Udorn/307/1972 (H3N2)	•••YV••D•G	K•YQ•A•G••	TT•DN••••D	••H••T•H••	•LMNEL•VPF	HL-GT•QVCI	
A/Shanghai/02/2013 (H7N9)	•••YV••D•D	•••FYA•S••	TTIRG •••••	••H•••QY•A	•I•W•LSSP•	TV····V·CI	
	181	_					
A/Texas/05/2009(H1N1)	AWSASACHDG	<b>IN</b> WLTIGISG	PDNGAVAVLK	YNGIITDTIK	SWRNNILRTQ	ESECACVNGS	
A/Vietnam/1203/2004 (H5N1)		TS·····			• • • • • • • • • • • •	•••••	
A/Udorn/307/1972 (H3N2)	•••s•s••••	KA••HVCVT•	Y•KN•T•SFI	•D•RLV•S•G	••SQ•••••	••••V•I••T	
A/Shanghai/02/2013 (H7N9)	G••STS••••	KSRMS•C•••	•N•N•S••VW	••RRPVAE•N	T•AR•••••	••••V•H••V	
	241						
A/Texas/05/2009(H1N1)	CFTVMTDGPS	NGQAS <mark>Y</mark> KIF <mark>R</mark>	IEKGKIVKSV	EMNAPNYHYE	ECSCYP <mark>DSS</mark> E	ITCVCRDNWH	
A/Vietnam/1203/2004 (H5N1)		••••H•••K	M••••V••••	• LD • • • • • •	•••••NAG•		
A/Udorn/307/1972 (H3N2)	•TV••••SA	S•R•DT••LF	••E••••HIS	PLSGSAQ•V•	•••••RYPG	VR•I••••K	
A/Shanghai/02/2013 (H7N9)	• PV• F••• SA	T•P•DTR•YY	FKE•••L•WE	SLTGTAK•I•	••••GERTG	••••T••••Q	
						344	
	301						
A/Texas/05/2009(HINI)	GSNRPWVSFN	-QNLEYQIGY	ICSGIFGDNP	RPNDKTGS	C-GPVSSNGA	NGVKGFSFKY	
A/Udorp/307/1972 (H3N2)	••••V•DT•	VKDYSIDSS.	V•••I.V••T•	•N••RSSNSY	• BN • NNEK • N	H••••WA•DD	
A/Shanghai/02/2013 (H7N9)	·····VIOID	PVAMTHTSO.	•••PVLT•••	••••PNIGK-	•ND•YPG•NN	·····YLD	
	3.61	~ 365,366					
A/Texas/05/2009(H1N1)	GNGVWIGBTK	SISSBNGEEM	TWDPNGWTGT	DNNESTKODT	VGIN-EWSGY	SGSEVOHPET	
A/Vietnam/1203/2004 (H5N1)	•••••	•TN••S••••	•••••E•	•SS••V••••	•A•T-D••••	•••••	
A/Udorn/307/1972 (H3N2)	••D••M•••I	•ED••S•Y•T	FKVIG••STP	NSKLQ•NRQV	IVDSDNR•••	••I•SV	
A/Shanghai/02/2013 (H7N9)	• ANT • L • • • I	•TA••S•Y••	LKV••AL•DD	RSK-P•QGQT	IVL•AD••••	••••MDYW-A	
	421						
A/Texas/05/2009(H1N1)	TGLDCIRPCF	WVELIRGRPK	ENTI-WTSGS	SISFCGVNSD	TVGWSWPDGA	ELPFTIDK	
A/Vietnam/1203/2004 (H5N1)	•••••	• • • • • • • • • • •	• S • • - • • • • •	• • • • • • • • • • •	•••••	•••••	
A/Udorn/307/1972 (H3N2)	E•KS••NR••	Y••••EQ	•TRVW•••N•	IVV · · · TSGT	YGTG•••••	DINLMPI-	
A/Shanghai/02/2013 (H7N9)	$E \bullet D - \bullet Y \bullet A \bullet \bullet$	Y • • • • • • • • • • • • • • • • • • •	• DKVW • • • N •	IV•M•SSTEF	LGQ•N••••	KIEYFL	

FIG 7 Sequence alignments on influenza A virus NA proteins. The full lengths of four influenza NA sequences, from A/Texas/05/2009 (pH1N1), A/Vietnam/1203/2004 (H5N1), A/Udorn/307/1972 (H3N2), and A/Shanghai/02/2013 (H7N9), were aligned using the Influenza Virus Resource (http://www.ncbi.nlm.nih.gov/genomes/FLU/FLU.html) and Vector NTI software. Identical amino acids are indicated by dots, and noncorresponding amino acid sites are shown by dashes. Identical amino acids (94.6%) comprising 34 different residues between the NA ectodomain of A/Texas/05/2009 (pH1N1) and A/Vietnam/1203/2004 (H5N1) are shown by red highlighting. The four target residues selected from those 34 (I149V, N344Y, I365T, and S366N) are highlighted in blue.

amino acid differences at residues 118 or 119 (on  $\beta_1L_{01}$ ), 151 or 152 (on  $\beta_1L_{23}$  [ $L_{23}$  denotes a loop between strands 2 and 3 within a sheet]); 198 ( $\beta_2L_{23}$ ), 224 or 227 ( $\beta_3L_{01}$ ), 243 ( $\beta_3L_{23}$ ), 274, 276, or 277 ( $\beta_4L_{01}$ ), 292 ( $\beta_1L_{23}$ ), 330 or 350 ( $\beta_5L_{01}$ ), or 425 ( $\beta_6S_2$ ) (11). However, we did find four amino acid differences between pH1N1 rNA and H5N1 rNA (I149V, N344Y, I365T, and S366N) close to enzymatic catalytic pockets consisting of the  $\beta_1L_{23}$  loop (151 and 152) and the  $\beta_5L_{01}$  loop (330 and 350).

NI antibodies are important for NA-based vaccine development. Other researchers have reported that NI antibodies induced by pH1N1 immunizations are cross-reactive with H5N1 viruses in mice (34) and ferrets (21, 22). Our results indicate that the NI antibodies induced by pH1N1 rNA immunizations (20  $\mu$ g), with an IC<sub>50</sub> of 2.78, resulted in a 60% survival rate after H5N1 virus challenges. In contrast, no protection against H5N1 virus challenges was noted following immunizations with 2  $\mu$ g of pH1N1 rNA, with an IC<sub>50</sub> of 2.47 (Fig. 3I, 5D, and 6D), while 100% protection against pH1N1 virus challenges was observed in mice immunized with either 2 or 20  $\mu$ g of H5N1 rNA (IC<sub>50</sub>s of 3.53 and 3.75, respectively) (Fig. 3I, 5C, and 6C). The NI antibodies induced by 2- or 20- $\mu$ g H5N1 rNA immunizations had IC<sub>50</sub>s of 1.92 and 2.30, respectively, against the H3N2 virus, and the NI antibodies induced by 2- or 20- $\mu$ g pH1N1 rNA immunizations had IC<sub>50</sub>s of 2.03 and 2.32, respectively (Fig. 3I). The NI antibodies induced by 2- or 20- $\mu$ g H5N1 rNA immunizations had IC<sub>50</sub>s of 1.45 and 2.26 against the H7N9 virus, respectively, and the NI antibodies induced by 2- or 20- $\mu$ g pH1N1 rNA immunizations had IC<sub>50</sub>s of 1.45 and 2.26 against the H7N9 virus, respectively, and the NI antibodies induced by 2- or 20- $\mu$ g pH1N1 rNA immunizations had IC<sub>50</sub>s of 1.31 and 2.17, respectively. In other words, H5N1



FIG 8 Three-dimensional neuraminidase structure diagram. The 3-D protein structures of the tetrameric H5N1 rNA proteins were modified using the crystal structure of NA from A/Vietnam/1203/04 strain (PDB 2HTY). The top and side views of the monomeric structures were diagramed by using PyMOL software, and the enzyme active sites are shown as red sticks. The four target residues (I149V, N344Y, and I365T/S366N) are shaded in blue.

rNA immunizations induced more potent cross-reactive NI antibodies and protective immunity against heterologous pH1N1 viruses compared to the pH1N1 rNA protective immunity against heterologous H5N1 viruses. These results are in agreement with those from studies of pH1N1 rNA (A/California/04/2009) immunization against an H5N1 (A/Turkey/1/2005) virus strain (NI antibody titer of 2.67) (19) and of NA antisera induced by live, attenuated pH1N1 (A/California/7/2009) that cross-reacted with various H5N1 strains in ferrets (NI antibody titers of  $\sim$ 2.2 to 2.3) (21). However, in neither case was any protection observed against the heterosubtypic H7N9 virus challenges (Fig. 3I, 5E, and 6E). Again, our findings are similar to those of a recent report using rNA immunizations that showed no protection against heterosubyptic influenza virus challenges (29).

Conserved influenza NA epitopes are of great interest to researchers working on a universal influenza vaccine. According to one report, 20 amino acid residues in 7 upper surface loops (195 to 202, 216 to 231, 243 to 251, 316 to 353, 364 to 374, 398 to 407, and 428 to 439) close to enzymatically active sites may be key to eliciting cross-reactive NI antibodies between H1N1 (A/Beijing/262/ 95) and H5N1 (A/Hong Kong/483/97) (35). Another research team reported that N1 conserved epitopes at NA residues 273, 338, and 339 induced cross-reactive NA-specific antibodies and provided protection against seasonal H1N1, 1918 H1N1, 2009 pandemic H1N1 (pH1N1), and H5N1 avian influenza viruses (34). To date, the only NA epitopes that are universally conserved among all influenza A viruses have been reported at NA residues 222 to 230 for the formation of active sites at residues 224 and 227  $(\beta_3 L_{01})$  (36). In the present study, we found four amino acid residues close to active enzyme sites in pH1N1 (A/Texas/05/2009) and H5N1 (A/Vietnam/1203/2004): I149V, N344Y, I365T, and S366N (Fig. 8). We also investigated three mutant pH1N1 rNA proteins (I149V, N344Y, and I365T/S366N mutants) by using mutagenesis to map these amino acids that may contribute to cross-reactive NI antibody elicitation. Our results indicate that the pH1N1 rNA (I365T/S366N) mutant protein was the most active in eliciting increased NI antibody titers against homologous, het-



FIG 9 Construction, expression, and characterization of WT and mutant pH1N1 rNAs. (A) The wild and mutant pH1N1 rNAs were constructed by a site-directed mutagenesis strategy and expressed using a baculovirus-insect cell expression system. (B and C) Purified rNA proteins (2 µg) were confirmed by Coomassie blue staining (B) and Western blotting (C).



FIG 10 NA-inhibiting-antibody curves and the corresponding  $IC_{50}$ s against different virus strains produced by immunizing mice with the WT and mutant pH1N1 rNAs. Fetuin-based assays were used to measure NA-inhibiting-antibody titers as reductions in the NA enzymatic activities of the pH1N1 or H3N2 viruses or H5N1 or H7N9 VLPs. (A to D) Inhibition curves for the WT pH1N1 rNA or mutant pH1N1 rNAs (I149V, N344Y, or I365T/S366N) against homologous pH1N1 viruses (A), the heterosubtypic H5N1 VLPs (B), the H3N2 viral strain (C), or H7N9 VLPs (D) are plotted. (E) The corresponding  $IC_{50}$ s were determined as the 50% reduction in the NA enzymatic activities of the homologous strains (white bars) or heterosubtypic viruses (filled bars). Integrated data are expressed as means  $\pm$  SD from three experiments. \*, P < 0.05; \*\*, P < 0.01; \*\*\*, P < 0.001.

erologous, and heterosubtypic viruses (Fig. 10). These findings provide useful information for the development of an NA-based universal influenza vaccine.

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