Broad Humoral and Cellular Immunity Elicited by a Bivalent DNA Vaccine Encoding HA and NP Genes from an H5N1 Virus

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Abstract

Influenza A virus is highly variable and a major viral respiratory pathogen that can cause severe illness in humans. Therefore it is important to induce a sufficient immune response specific to current strains and to heterosubtypic viruses with vaccines. In this study, we developed a dual-promoter-based bivalent DNA vaccine that encodes both hemagglutinin (HA) and nucleoprotein (NP) proteins from a highly pathogenic A/Chicken/Henan/12/2004 (H5N1) virus. Our results show that the expression levels of HA and NP genes from the dual-promoter plasmid are similar to those seen when they are expressed individually in independent plasmids. When the bivalent DNA vaccine was inoculated via intramuscular injection and *in vivo* electroporation, high levels of both humoral and cellular immune responses were elicited against homologous H5N1 virus and heterosubtypic H9N2 virus. Furthermore, no obvious antigenic competition was observed between HA and NP proteins in the dual-promoter-based bivalent vaccine compared to monovalent vaccines. Our data suggest that a combination of influenza surface and internal viral genes in a dual-promoter-expressing plasmid may provide a new approach for developing a DNA vaccine that may protect not only specifically against a currently circulating strain, but also may cross-protect broadly against new heterosubtypic viruses.

Introduction

THE AVIAN INFLUENZA VIRUSES WERE THOUGHT UNABLE to be transmitted directly from birds to humans until 1997, when a highly pathogenic avian influenza (HPAI) A virus H5N1 broke out in Hong Kong (12,16,40). Data from the World Health Organization (WHO) indicated 500 confirmed human cases of H5N1 virus infection, and the fatality rate for the virus is approximately 60% (50). In addition to the H5N1 virus, other avian influenza virus subtypes have also been reported to cause human infections (4,31,54). Although the slaughter of poultry efficiently eliminated the source of the infection, a human pandemic caused by an avian virus is still possible (6). Furthermore, potential reassortment between human flu viruses and avian flu viruses has caused global concern about the possibility of creating a new virulent strain that is more easily transmissible and lethal to humans (36). Therefore, vaccines that protect against infection or limit the

spread of avian influenza viruses, particularly HPAI viruses, are urgently needed.

Influenza A virus has been successful in its evolution due to its antigenic variation, which appears in two forms: antigenic shift and antigenic drift. This epidemiological property may cause the emergence of new epidemic and pandemic viruses annually. As a result, every year before the ensuing influenza season, the WHO recommends the composition of an influenza vaccine suitable for the upcoming year based on global surveillance. However, the seasonal vaccine reconstituted with the WHO-recommended strains may not always be capable of providing protection against a new emerging epidemic strain. The 2009 swine-origin influenza virus (S-OIV) H1N1 is such a case, and it broke out in April 2009 in Mexico and the U.S. (7,23,51). Analysis of its antigenic and genetic characteristics showed that this virus was a new reassortant virus introduced in humans, against which the population had little immunity (25,52). Vaccines against

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this novel virus were urgently produced worldwide to prevent a potential epidemic in the winter season. It is consequently believed that an influenza vaccine candidate that provides broad cross-strain protection is a promising way to deal with such situations in the future, and may provide protection against this ever-changing virus (18).

Thus an ideal vaccine against influenza virus must be not only specific to the currently circulating strain, but must also cover different subtypes. The most widely used influenza vaccines, the inactivated vaccines, fail to fulfill this criterion, as they are only able to elicit antibody responses against viral proteins, and thus are only effective for circulating strains (34). The live attenuated vaccines, including cold adapted vaccines and genetically engineered vaccines, appear to be ideal, as they can induce both antibody responses against circulating strains, and a cell-mediated immune response against internal viral proteins conserved among subtypes. However, a potential reassortment with natural virus increases the risk of this approach (34). In addition, the handling of live viruses needed to make these vaccines restrict its large-scale production. Thus the developing DNA vaccine approach is promising due to induction of both humoral and cellular immune responses, and its safety and low cost of production (19,21).

As a model system for DNA vaccines, influenza DNA vaccines have been proven in different animal models to provide protective immunity against viral infection (44). DNA constructs expressing the viral surface glycoproteins hemagglutinin (HA) and neuraminidase (NA) were demonstrated to provide efficient protection against homologous or drift viral infection, but little cross-subtype protection was elicited (10,29,30,37,45). Plasmids encoding internal conserved viral proteins, for example nucleoprotein (NP) and matrix protein 1 (M1), are able to elicit protection against heterosubtypic virus challenge via a cell-mediated immune response (19,20,46,47). It is therefore reasonable to propose that a combination of the surface and internal antigens of influenza virus may provide effective protection, not only against the circulating strain, but also broadly against heterosubtypic viruses.

In this study, we generated an effective bivalent DNA vaccine by encoding both HA and NP proteins from the HPAI virus A/Chicken/Henan/12/2004 (H5N1). The results show that this bivalent vaccine can elicit humoral and cell-mediated immune responses against both homologous and heterosubtypic viruses when administered via intramuscular (IM) injection and *in vivo* electroporation. Our work provides a new DNA vaccine candidate that is not only specific enough to neutralize HPAI H5N1 virus, but is also broadly responsive to heterosubtypic viruses.

Materials and Methods

Mice and viruses

Female BALB/c mice 4–6 weeks old were purchased from Shanghai SLAC Laboratory Animal Co., Ltd., and randomly divided into groups of 10 mice per group. All mice were maintained with free access to sterile food and water.

The A/Chicken/Henan/12/2004 (H5N1) virus inactivated with formaldehyde was kindly provided by Prof. Ze Chen (Shanghai Institute of Biological Products). A/Chicken/Jiangsu/7/2002 (H9N2) virus growing in Madin-

Darby canine kidney cells was collected and inactivated with formaldehyde.

Plasmid constructs for immunization

Viral cDNA from A/Chicken/Henan/12/2004 (H5N1) was kindly provided by Prof. Ze Chen. The full-length HA gene was cloned into the pBudCE4.1 vector (Invitrogen, Carlsbad, CA), under the control of the human elongation factor 1 α -subunit (EF-1 α) promoter, using the restriction enzyme NotI/XhoI (Takara Bio Inc., Shiga, Japan), and was designated pHA. The full-length NP gene was cloned into the pBudCE4.1 vector under the control of the human cytomegalovirus (CMV) immediate-early promoter using restriction enzyme BamHI/HindIII (Takara Bio Inc.), and was designated pNP. The full-length NP gene was further cloned into pHA under the CMV promoter to generate pHA/NP.

To produce H5N1 pseudotyped virus, the NA gene from A/Chicken/Henan/12/2004 (H5N1) was cloned into pHA under the CMV promoter, and was designated pHA/NA. Plasmids were grown in DH-5 α *E. coli* strain, and purified using commercial purification kits. All plasmids were confirmed by sequence analysis.

Cells and transfection

Human embryonic kidney 293T cells (HEK 293T; ATCC), and Madin-Darby canine kidney cells (MDCK; ATCC) were cultured in Dulbecco's modified Eagle's medium (DMEM) (Gibco, Carlsbad, CA) containing 10% fetal bovine serum (Gibco) at 37°C in a CO₂ incubator.

Lipofectamine (Invitrogen) was applied for transient transfection following the manufacturer's protocol.

Western blot analysis

Cells were lysed directly in sodium dodecyl sulfate (SDS) sample buffer (60 mM Tris-HCl [pH 6.8], 2% SDS, 10% glycerol, 5% 2-mercaptoethanol, and 0.01% bromophenol blue), followed by boiling for 10 min. Whole-cell lysates were further subjected to SDS-PAGE. The proteins were transferred to nitrocellulose membranes (BioRad Laboratories, Inc., Hercules, CA). The HA protein was detected with anti-HA polyclonal antibody (Antibody Research Centre, Shanghai Institute for Biological Sciences). NP protein was detected with anti-NP polyclonal antibody (Antibody Research Centre, Shanghai Institute for Biological Sciences), and actin was detected with rabbit anti-actin polyclonal antibody (Sigma-Aldrich, St. Louis, MO). Goat anti-rabbit IgG HRP-conjugated antibody (Southern Biotechnology Associates, Birmingham, AL) was applied as the secondary antibody. The exposure was developed with an enhanced chemiluminescence kit (Pierce Protein Research Products, Rockford, IL).

Immunization by *in vivo* electroporation

Plasmid DNA encoding viral proteins was inoculated by *in vivo* electroporation, which was carried out according to the method previously described (1). Mice (aged 4–6 wk) were immunized thrice (at 3-wk intervals) by injection into the quadriceps muscles of two legs with the same molar amount of the vector (pBudCE4.1), pHA, pNP, or pHA/NP, at doses of 50, 65, 63, and 83 μ g per mouse, respectively. A

pair of 5-mm-separated electrode needles was then inserted into the muscle to cover the DNA injection sites, and electric pulses were delivered using an electric pulse generator (Electro Square Porator T830 M; BTX, San Diego, CA) (8,15). *In vivo* electroporation parameters were: 100 V, 5 pulses, 20 msec pulse duration, and 200 msec between pulses.

Detection of anti-influenza-virus antibody

Antibody levels against H5N1 (A/Chicken/Henan/12/2004) and H9N2 (A/Chicken/Jiangsu/7/2002) were assayed by enzyme-linked immunosorbent assay (ELISA). In brief, 96-well microtiter plates were coated with 5 µg/mL of chemically-killed corresponding virus at 4°C overnight. Following the blocking of each well with 10% bovine serum-PBST at 37°C for 2 h, the mouse sera collected on the indicated days were prepared in serial dilutions, and were added to each well and incubated at 37°C for 2 h. A secondary goat anti-mouse IgG HRP-conjugated antibody (Sigma-Aldrich) at 1:1000 dilution was added to each well and incubated at 37°C for 1 h. To develop the ELISA, a 10-mg TMB tablet (Sigma-Aldrich) was dissolved in 1 mL of 0.025 M phosphate-citrate buffer, and 50 µL of the resulting solution was added to each well. The reaction was stopped by the addition of 0.2 M H₂SO₄, and the resulting optical density (OD) at 450 nm was analyzed with a plate reader.

Production of H5N1 pseudotyped virus and neutralization assay

Pseudotyped virus was produced to avoid the manipulation of live HPAI virus in the neutralization assay as previously described (3,11). Briefly, HEK293T cells were plated at approximately 50% confluence in six-well plates. Twenty-four hours later, the plasmid pHA/NA encoding the glycoproteins HA and NA from A/Chicken/Henan/12/2004 (H5N1) virus was co-transfected with packaging plasmid pNL4-3^{Luc+Env-Vpr-} to generate a pseudotyped HIV-backbone luciferase-reporter virus with influenza glycoproteins embedded on the surface. Supernatants from the HEK293T cells were collected at 48 h post-transfection and aliquotted after clarification by low-speed centrifugation (3000 rpm for 5 min), and kept frozen.

For the neutralization assay, MDCK cells were plated as target cells in 24-well plates. During infection, 100 µL H5N1 pseudotyped virus was incubated with twofold serial dilutions of antisera (starting dilution 1:500) in 900 µL DMEM at 37°C for 1 h. Subsequently, the medium was aspirated and 1 mL virus-antiserum mixture was transferred onto the cells. Twenty-four hours later, 1 mL fresh DMEM without FBS was added, and the cells were lysed with passive lysis buffer (Promega Corp., Madison, WI) 48 h later. Because the pseudotyped virus is env⁻, it is competent for only a single cycle of virus replication. Thus the intracellular amount of luciferase activity represents a direct reflection of virus entry. Luciferase assay was performed with Steady-Glow Luciferase Substrate (Promega Corp.).

Splenocyte proliferation activity assay

On the seventh day after the third injection, the mice were sacrificed and a single-cell suspension was prepared from the spleens of each group. The splenocytes were cultured in

triplicate in 96-well plates at 4×10^5 cells per well in RDF medium (Gibco) with 10% fetal bovine serum (Gibco). The cultures were then stimulated for 36 h with each of the following: 10 µg/mL BSA (irrelevant antigen), 2 µg/mL Con A (positive control), and 10 µg/mL of the chemically-killed H5N1 virus, or 10 µg/mL of the chemically-killed H9N2 virus. The cells were then incubated at 37°C and 5% CO₂ for 36 h, then 1 µCi ³[H]thymidine (Amersham Pharmacia Biotech, Piscataway, NJ) was added to each well for 8–12 h. The cells were harvested and ³[H]thymidine uptake (cpm) was counted on a Beckman LS 6500 scintillation counter (Beckman Coulter, Fullerton, CA). The stimulation indexes (SI) were determined by cpm reading of antigen-containing wells divided by cpm reading of wells without antigen.

In vivo CTL assay

The *in vivo* cytotoxicity assay was performed as previously documented (13,33). To prepare target cells, splenocytes were obtained from naïve BALB/c spleen cell suspensions after erythrocytes were removed. The cells were then washed and split into two populations. One population was pulsed with 10⁻⁶ M of a 1:1 peptide mixture (HA:YSTVASSL or NP:TYQRTALV) (17), incubated at 37°C in 5% CO₂ for 1 h, washed, labeled with a high concentration of CFSE (carboxyfluorescein diacetate *N*-succinimidyl ester; Sigma-Aldrich) (5 µM), and designated CFSE^{high} cells. The other population was left without peptide, labeled with a low concentration of CFSE (0.5 µM), and designated CFSE^{low} cells. For *in vivo* injection, a mixture of equal numbers of cells from the CFSE^{high} and CFSE^{low} populations (2×10^7 cells in 400 µL of PBS in total) was injected intravenously into each immunized mouse on the seventh day after the third immunization. Four hours after injection, the mice were sacrificed for their splenocytes. Cell suspensions were analyzed on a FACSCalibur analyzer (BD Biosciences Pharmingen, Franklin Lakes, NJ), and each population was detected by its differential CFSE fluorescence intensity. Up to 1×10^4 CFSE-positive cells were collected for analysis. To calculate specific lysis, the following formulas were used: ratio = (percentage CFSE^{low}/percentage CFSE^{high}); percentage specific lysis = $[1 \times (\text{ratio unprimed}/\text{ratio primed}) \times 100]$ (27).

Cytokine assay

On the seventh day after the third immunization, splenocytes were harvested for antigen restimulation. First, 4×10^5 cells in 96-well plates were stimulated in culture with 10 µg/mL BSA (irrelevant antigen), 2 µg/mL ConA (positive control), 10 µg/mL chemically-killed H5N1 virus, or 10 µg/mL chemically-killed H9N2 virus, at 37°C in 5% CO₂ for 72 h. The levels of secreted IFN-γ and IL-4 in the cultured supernatant were determined using ELISA kits from R&D Systems (Minneapolis, MN).

Data analysis

All values were expressed as mean ± standard deviation (SD). Statistical analysis of the experimental data was performed using the two-tailed Student's *t*-test or one-way analysis of variance (ANOVA). For all tests, *p* < 0.05 was considered statistically significant.

Results

Expression of DNA constructs in eukaryotic cells

To test whether a bivalent DNA vaccine containing HA and NP from an HPAI H5N1 virus could induce both strong cellular and humoral immunity, the entire length of the HA and NP coding regions from A/Chicken/Henan/12/2004 (H5N1) virus were cloned into a dual-promoter eukaryotic-expressing vector pBudCE4.1 under their respective promoters as shown in Fig. 1A. The pHA/NP plasmid expressing two viral proteins at the same time was used as the bivalent DNA vaccine, while the monovalent plasmids pHA and pNP, each encoding only one viral protein, and vector plasmid pBudCE4.1, were included as controls. To confirm the expression of all the DNA constructs, human 293T cells were transfected with same molar amount of the corresponding plasmid, and protein expression was analyzed by Western blot assay with anti-HA or anti-NP polyclonal antibody. As shown in Fig. 1B, the co-expression of HA and NP in pHA/NP was similar to the levels of pHA or pNP expressed singly. These data suggest that the bivalent DNA vaccine was able to express successfully and efficiently in eukaryotic cells.

HA expression triggers a predominantly humoral immune response with neutralizing activity, and NP elicits a cross-response to heterosubtypic virus

To examine the humoral response induced by the bivalent DNA vaccine, we first examined the level of anti-H5N1-

specific antibodies in mice immunized with the respective DNA constructs. Sera of vaccinated mice from days 21, 35, and 49 post-initial immunization were assayed by ELISA, and the titers of anti-H5N1-specific IgG were analyzed. The data in Fig. 2A show that mice immunized with pHA, pNP, and pHA/NP induced significantly high levels of specific antibodies against chemically-killed H5N1 virus, compared to the vector-immunized group at day 49. Moreover, the highest antibody titer was observed in the group immunized with the bivalent DNA plasmid pHA/NP, which was proposed to elicit both anti-HA and anti-NP antibodies.

As neutralizing antibody is responsible for efficiently preventing virus infection, we further examined the neutralizing activity of the antisera. To this end, H5N1-pseudotyped virus was obtained by co-expression of the viral HA and NA genes on the surface of a HIV-based packaging virion. The virus-neutralizing abilities of the sera from all groups were assessed using reductions in luciferase expression of the H5N1-pseudotype virus (3). The results clearly showed that of all immunized groups, only the pHA and pHA/NP groups were able to induce a strong neutralizing-antibody response (Fig. 2B). In contrast, little neutralizing antibody was elicited in the group immunized with pNP or vector, indicating that the HA protein of influenza virus is the predominant inducer of neutralizing antibody.

To test whether our bivalent DNA vaccine had broad response against heterosubtypic virus, we then examined the heterosubtypic IgG against avian influenza subtype H9N2 virus, which is also able to cause cross-species infection in hu-

FIG. 1. Construction and eukaryotic expression of DNA vaccines. (A) Schematic diagram of DNA vaccine constructs. The full-length HA and NP genes of H5N1-HPAIV strain A/Chicken/Henan/12/2004 was cloned into the dual-promoter plasmid pBudCE4.1 for eukaryotic expression. (B) Expression of DNA constructs in HEK293T cells. The HEK293T cells were transfected with equal molar amounts of the indicated plasmids, and cell lysates were collected 24 h post-transfection and subjected to Western blot analysis with anti-HA and anti-NP polyclonal antibodies. The β -actin was detected with anti-actin-specific polyclonal antibody as a loading control.

FIG. 2. Generation of anti-H5N1 homologous and anti-H9N2 heterosubtypic IgG in mice. (A) Antibody response to A/Chicken/Henan/12/2004 (H5N1) virus. Mice were bled on days 21, 35, and 49 after the initial immunization, and sera were prepared in serial dilutions. The antibodies against the H5N1 virus were evaluated by ELISA with 5 μ g/mL of chemically-killed A/Chicken/Henan/12/2004 (H5N1) virus as antigen coated on each well in a 96-well plate. The titers of the sera were defined as the last serial dilution with an OD₄₅₀ value exceeding the cut-off value (cut-off = 2 times the OD value of naïve serum), and were represented as the mean titers + standard deviation in five animals (* p < 0.05 by Student's *t*-test compared to the vector-immunized group). (B) Neutralizing activity of sera from immunized mice. Immunized sera were collected on day 49 after initial immunization, and prepared in serial dilutions. Neutralizing activities of the sera were determined by measuring the inhibition of H5N1 pseudotyped virus infectivity on MDCK cells (–, negative control in which no pseudotyped virus was added to the cells; +, positive control in which pseudotyped virus mixed with irrelevant antibody was added to the cells, defined as 100% infectivity; * p < 0.05 compared to the vector- and pNP-immunized groups). (C) Antibody response to A/Chicken/Jiangsu/7/2002 (H9N2) virus. The antibody against the H9N2 virus was evaluated by ELISA with 5 μ g/mL of chemically-killed A/Chicken/Jiangsu/7/2002 (H9N2) virus antigen coated on each well in a 96-well plate. The titers were calculated as above (* p < 0.05 compared to the vector- and pHA-immunized groups).

mans. Antibody titers in sera from immunized mice were determined using ELISA against chemically-killed A/Chicken/Jiangsu/7/2002 (H9N2) virus. As shown in Fig. 2C, the bivalent pHA/NP and the monovalent pNP immunizations elicited similar amounts of high-level anti-H9N2-specific antibodies, while antibody titers in both the vector- and pHA-immunized groups were quite low.

Taken together, these data suggest that only the bivalent DNA vaccine pHA/NP was able to generate both anti-H5N1 HA-specific neutralizing antibody and anti-H9N2 NP-heterosubtypic antibody.

NP expression induces stronger splenocyte proliferation with a heterosubtypic response

To determine the splenocyte response, a single suspension of splenocytes was prepared from mouse spleens on the seventh day after the third injection and assayed. As shown in Fig. 3, when stimulated with chemically-killed A/Chicken/Henan/12/2004 (H5N1) virus, cells isolated from the spleens of mice injected with both pNP and pHA/NP achieved high stimulation indexes. The splenocyte proliferation in mice injected with pHA was also observable, but was relatively lower compared to the DNA constructs expressing NP protein (pNP and pHA/NP), while the highest proliferation was observed in mice injected with pHA/NP. When stimulated with chemically-killed heterosubtypic A/Chicken/Jiangsu/7/2002 (H9N2) virus, significant splenocyte proliferation was observed in mice injected with DNA

constructs pNP or pHA/NP, but not in the pHA group. These data in all indicate that immunization with DNA constructs expressing NP protein (pNP and pHA/NP) elicit more efficient and robust splenocyte proliferation responses against both homologous H5N1 and heterosubtypic H9N2 viruses.

The strongest CTL response was elicited by the bivalent pHA/NP immunization

Virus-specific cytotoxic T lymphocyte (CTL) activity has been reported to be important in controlling virus infections via direct cytolysis of virus-infected cells. We therefore set out to determine the CTL activity using an *in vivo* CTL assay, which reflects more of the true stimulation than the *in vitro* assay. To detect the presence of anti-influenza virus-specific CTL *in vivo*, we adopted a previously documented method (13,33). A mixture (1:1) of influenza virus major histocompatibility complex class I (MHC-I) peptides (IYSTVASSL from the HA protein and LYEKVKSQ from the NP protein), which are mostly conserved among influenza strains and subtypes, was applied to stimulate targeted splenocytes labeled with a high concentration of CFSE (CFSE^{high}). The same number of untargeted splenocytes labeled with a low concentration of CFSE (CFSE^{low}) was injected together with the CFSE^{high} cells into immunized mice intravenously. The CTL response was then evaluated, as peptide-labeled cells were killed, while antigen-unlabeled cells survived. The results in Fig. 4A show that all of the pHA, pNP, and

FIG. 3. T-cell proliferation after DNA vaccination. A single suspension of splenocytes of mice immunized with the indicated DNA constructs was isolated on the seventh day after the third immunization, and stimulated *in vitro* either with chemically-killed A/Chicken/Henan/12/2004 (H5N1) virus, with A/Chicken/Jiangsu/7/2002 (H9N2) virus, with an irrelevant protein (BSA) as a negative control, or with ConA as a positive control. The stimulation indexes ($\pm$ standard deviation) were derived as the value of the test sample divided by the value of blank medium (* $p < 0.05$, ** $p < 0.01$ compared to the BSA-stimulated index within each group; * $p < 0.05$, ** $p < 0.01$ indicate statistical significance when the indicated columns were compared between groups).

FIG. 4. *In vivo* influenza-virus-specific CTL assay. A 1:1 mixture of 10^7 cells of each target cell population was injected intravenously into mice 7 d after the third injection with the indicated DNA vaccines. After 4 h, the mice were killed and the splenocytes were analyzed for the presence of CFSE^{high} (pulsed with influenza epitopes), and CFSE^{low} (unpulsed) target cell populations. (A) Representative histogram plot of splenocytes obtained from a mouse vaccinated with each DNA construct at 4 h post-transfer of CFSE-labeled target cells. (B) The percentage of specific lysis (+standard deviation) in five animals was calculated as described in the materials and methods section (* $p < 0.05$, ** $p < 0.01$ compared to the vector-immunized group; * $p < 0.05$ for the pHA/NP-immunized group compared to the pHA-immunized group).

pHA/NP immunizations induced significantly efficient lysis of influenza peptide-labeled cells, compared to that seen in the vector-immunized group. The quantitative analysis in Fig. 4B further suggests that pNP immunization induced higher CTL activity than pHA immunization, while the

highest CTL activity was observed in mice vaccinated with the bivalent pHA/NP construct. These data in all indicate that the strongest CTL response was elicited by the bivalent DNA vaccine pHA/NP, compared to the monovalent constructs.

Both IL-4 and IFN- γ are induced by the bivalent pHA/NP immunization

The ability of this bivalent DNA vaccine (pHA/NP) to elicit both humoral and cellular immune responses was examined by analyzing its effect on the Th-1- and Th-2-stimulated cytokine response. ELISA was used to measure the levels of IFN- γ , representing the Th-1 cytokine, and IL-4, representing the Th-2 cytokine, in restimulated splenocytes taken from mice on the seventh day after the third immunization. Chemically-killed A/Chicken/Henan/12/2004 (H5N1) and A/Chicken/Jiangsu/7/2002 (H9N2) viruses were applied as homologous

and heterosubtypic antigens, respectively; mitogen concanavalin A (ConA) was used as a positive control; and BSA was used as an irrelevant control. As shown in Fig. 5, when re-stimulated with homologous H5N1 virus, the pHA, pNP, and pHA/NP immunizations induced high levels of IFN- γ and IL-4 secretion, compared to vector immunization. However, only the DNA constructs expressing NP protein were able to elicit a high level of IFN- γ and IL-4 against heterosubtypic H9N2 virus. In all, the cytokine profile seen after DNA immunizations indicates that the bivalent DNA vaccine pHA/NP elicits high levels of both IFN- γ and IL-4 against both homologous and heterosubtypic virus.

FIG. 5. Effect of DNA vaccinations on IFN- γ and IL-4 release. Splenocytes harvested on the seventh day after the third immunization were stimulated *in vitro*, with chemically-killed A/Chicken/Henan/12/2004 (H5N1) virus, A/Chicken/Jiangsu/7/2002 (H9N2) virus, with an irrelevant protein (BSA) as a negative control, or with ConA as a positive control. Cytokine ELISA detection kits were used to evaluate IFN- γ (A) and IL-4 (B) levels. The concentrations of the released cytokines ($\pm$ standard deviation) were calculated using a cytokine standard curve on the same measuring plate (* $p < 0.05$ by Student's *t*-test compared to the BSA-stimulated value).

Discussion

Of the vaccine types against influenza virus, DNA vaccines are among the most promising vaccines, as they are capable of inducing both humoral and cellular immune responses. In this study, we developed a bivalent DNA vaccine encoding both the HA and NP proteins from a HPAI H5N1 virus. The combination of strain-specific surface viral protein, and conserved internal viral protein, makes the bivalent DNA vaccine capable of responding against both homologous and heterosubtypic viruses. This may be a potential vaccine candidate for use against both HPAI H5N1 viruses and any unexpected newly-emerging strains.

To design a DNA vaccine against influenza A virus, the surface viral protein HA is widely considered to be the predominant immunogen for the induction of strain-specific antibodies to prevent infection. However, with its many variations, the HA region alone can provide only weak broadly-protective immune activity (26), while the internal viral proteins, which are less variable, are reported to provide cross-subtype CTL-mediated protection (24,46). Therefore, a combination of surface and internal proteins should be both effective against the current circulating strain, and protective against other antigenic variants. The first evidence for this idea was seen with successful protection from antigenically-distinct influenza isolates in ferrets immunized with a combination of independent DNA plasmids encoding HA and two internal conserved proteins (19,20). However, few studies have followed this combination approach, possibly due to concerns about antigenic competition among viral proteins. Although a previous study showed no antigenic competition among the dissociated NA, M, and NP proteins of influenza A virus (28), no such data existed about the HA protein and internal viral proteins, when given either with independent plasmids or with one plasmid expressing multiple viral proteins. Our study with the bivalent DNA vaccine pHA/NP, expressing two viral proteins in a dual-promoter plasmid, demonstrates that there is no obvious antigenic competition between HA and NP. In most of the immune responses we have examined, bivalent DNA vaccines have exhibited relatively higher activity than any of the monovalent vaccines against the homologous H5N1 virus. Moreover, the similar immune responses seen with the bivalent pHA/NP and the univalent pNP vaccines against heterosubtypic H9N2 virus also indicates that antigenic competition between HA and NP proteins is minimal. As a consequence, a combination of surface HA and internal NP genes in a DNA vaccine is able to elicit both strain-specific and cross-subtype immune responses without inducing antigenic competition.

Among the delivery methods of DNA vaccination, the highest efficacy was achieved by *in vivo* electroporation and gene gun delivery (49). Electroporation of DNA vaccines *in vivo* has been proven to be an effective method to increase antigen expression and to improve immune responses in many animal models (2,38,42,53). *In vivo* electroporation has also been reported to successfully deliver influenza DNA vaccine against HPAI H5N1 virus in mice (9,58). In this study, *in vivo* electroporation was used to deliver the bivalent DNA vaccine with optimized parameters (8). Our results showed that high levels of both humoral and cellular immune responses were induced by the electroporation of bivalent

DNA vaccine without any adjuvant. Therefore, results from both previous and our studies provide promising evidence that electroporation appears to overcome the cellular uptake barrier of DNA vaccines.

To evaluate the neutralizing antibody response induced by our bivalent DNA vaccine, a pseudotyped virus-based assay was developed to avoid the need for the use of highly pathogenic H5N1 live virus. The conventional methods used to estimate neutralizing antibody responses against influenza viruses include the microneutralization (MN) and hemagglutination inhibition (HI) assays. However, both the MN and HI assays have their shortcomings. The MN assay requires the use of highly-pathogenic infectious virus, and therefore requires biosafety level 3 facilities, and is thus labor-intensive. Moreover, when evaluating the cytopathic effects, the accuracy of the MN assay is prone to the subjectivity of the researcher. With the HI assay, although it is easier to perform, differences in erythrocyte origins will cause differing HI titers. The lentiviral pseudotyped virus has been applied to evaluate the neutralizing antibody response against HPAI viruses (32,43,56). As the pseudotyped virus undergoes a single round of infection and cannot produce progeny viruses, highly-biosafe facilities are not required. Further, the pseudotyped virus contains the luciferase reporter gene, and infectivity can thus easily be measured by reading luminescence values. Moreover, the neutralizing antibody response evaluated using pseudotyped virus-based assays have also been found to be comparable to or even more sensitive than the MN and HI assays (43). In this study, the pseudotyped virus was used to express the HA and NA proteins from the same H5N1 strain to design a DNA vaccine; we found that both pHA and pHA/NP vaccinations elicited high levels of neutralizing antibody. This result indicates that DNA vaccines expressing the influenza HA protein are able to induce neutralizing antibodies with high efficacy.

To evaluate the heterosubtypic immune response induced by the bivalent DNA vaccine, we examined both the anti-heterosubtypic humoral antibody response, and the cell-mediated splenocyte proliferation and CTL response. For detection of anti-heterosubtypic antibody, chemically-killed A/Chicken/Jiangsu/7/2002 (H9N2) virus was used as antigen, and a high level of anti-NP antibody was observed, indicating an efficient cross-subtype humoral immune response. A previous study indicated that recombinant NP vaccination was able to generate protective anti-NP antibodies that reduced influenza virus titers and reduced mortality in mice (5). Although NP antibody may not protect cells against influenza infection (39), the substantial antibody response generated against NP protein by natural infection or vaccination indicates a potential role for anti-NP antibody in disease control and virus clearance (14). Considering that NP is also expressed on the surface of influenza-infected cells, and is a major target for CTL (35,39,48), it is reasonable to propose a significant role for the NP antibody in a cross-subtype immune response, perhaps even in humans. For the *in vivo* CTL assay, conserved HA and NP epitope-pulsed splenocytes were used as target cells to represent a general CTL response against influenza A virus. Compared to pHA, vaccination with either pNP and pHA/NP induced stronger cell-mediated immune responses, including splenocyte proliferation and a CTL response against both types of viruses.

These results are consistent with those of previous studies, indicating that influenza's internal proteins, particularly NP protein, are major targets for cell-mediated immunity (57). Both the anti-NP antibody and NP-induced cell-mediated immune responses may provide cross-reactive anti-influenza A virus protection.

The production of vaccines against HPAI viruses poses safety problems with regard to vaccine manufacture (55). The HPAI virus, like the avian H5N1 virus, kills chicken embryos in short order, resulting in substantial reductions in vaccine yield and potential contamination by the dead eggs. More seriously, production staff would be immunologically naïve to an avian virus, and thus would be susceptible to infection. Although the reverse-genetics technique based on plasmid rescue of infectious virions could create non-pathogenic vaccine strains by excising the extra basic amino acids at the cleavage site of the avian HA segment (41), relatively low efficiency in recombinant virus growth and the large demand for chicken embryos remain problematic in maintaining the global vaccine supply. Plasmid DNA vaccines are capable of overcoming all of the above problems, allowing for easier manipulation and faster production compared to traditional influenza vaccines. Several DNA vaccines against H5N1 virus have been described, using either HA-expressing plasmid, or internal M- and NP-expressing plasmids (9,22,29), though few studies have combined the surface and internal viral proteins to study the hetero-subtypic immune response. In this study, we constructed a bivalent DNA vaccine simultaneously expressing HA and NP proteins from an HPAI H5N1 virus. The surface HA protein expressed from the bivalent DNA vaccine induces an anti-H5N1-specific neutralizing antibody that can prevent viral infection, while the internal conserved NP protein expressed from the bivalent DNA vaccine induces both homologous and heterosubtypic anti-NP antibody, and cell-mediated immune responses that can aid in virus clearance. One can thus predict that the combined expression of influenza HA and NP by a bivalent DNA vaccine may be a new strategy to achieve protection against homo-subtypic influenza virus infection, as well as cross-protection against hetero-subtypic infections. This bivalent DNA vaccine may be a new vaccine candidate for use in the event of an HPAI pandemic.

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Author Disclosure Statement

No competing financial interests exist.

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