

Hemagglutinin-based polyanhydride nanovaccines against H5N1 influenza elicit protective virus neutralizing titers and cell-mediated immunity

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Abstract: H5N1 avian influenza is a significant global concern with the potential to become the next pandemic threat. Recombinant subunit vaccines are an attractive alternative for pandemic vaccines compared to traditional vaccine technologies. In particular, polyanhydride nanoparticles encapsulating subunit proteins have been shown to enhance humoral and cell-mediated immunity and provide protection upon lethal challenge. In this work, a recombinant H5 hemagglutinin trimer (H5₃) was produced and encapsulated into polyanhydride nanoparticles. The studies performed indicated that the recombinant H5₃ antigen was a robust immunogen. Immunizing mice with H5₃ encapsulated into polyanhydride nanoparticles induced high neutralizing antibody titers and enhanced CD4⁺ T cell recall responses in mice. Finally, the H5₃-based polyanhydride nanovaccine induced protective immunity against a low-pathogenic H5N1 viral challenge. Informatics analyses indicated that mice receiving the nanovaccine formulations and subsequently challenged with virus were similar to naïve mice that were not challenged. The current studies provide a basis to further exploit the advantages of polyanhydride nanovaccines in pandemic scenarios.

Keywords: polymer, nanoparticle, vaccine, subunit, neutralizing antibody

Introduction

Influenza virus is a major cause of serious respiratory illness and has been responsible for significant morbidity and mortality in humans worldwide. The major strains of influenza A virus circulating in human populations are H3N2 and H1N1, which are associated with seasonal influenza viral infections. In addition, the avian strains H5N1 and H7N9 have been found to infect humans;¹ however, they are not currently capable of sustained human-to-human transmission. Should these highly pathogenic avian viruses develop such capability, the disease could spread rapidly, resulting in a global influenza pandemic. Vaccination represents a critical control measure against yearly seasonal influenza viruses and is an essential component of pandemic preparedness plans.² Nearly all of the current influenza vaccine technologies are based on the use of embryonated eggs and require a relatively long production cycle, resulting in limited manufacturing capacity.² The response to the 2009 H1N1 pandemic clearly demonstrated the limitations of vaccine production methods with respect to rapid deployment, which led to vaccine shortages. Consequently, there is great interest in developing new technologies for rapid, large-scale production of safe and efficacious influenza vaccines.

Recombinant hemagglutinin (rHA)-based vaccines produced in mammalian or insect cell culture systems are attractive alternatives to egg-based vaccine technologies.³

Production and purification of rHA protein can effectively reduce the vaccine production time and the use of cell culture systems may glycosylate proteins, which is important for the development of neutralizing antibodies against receptor-binding site epitopes.⁴ Recent studies demonstrated that rHA engineered to form a stable trimeric configuration elicited a protective immune response in vaccinated animals.^{5–10} Neutralizing antibody levels and protection from disease were enhanced in mice vaccinated with soluble rHA trimers as compared to animals vaccinated with rHA monomers,^{5,8} indicating the importance of immunogens that mimic those expressed by the infectious agent. Along these lines, the composition and/or number of N-linked glycans on rHA trimers have been shown to modify the level of the protective antibody response.^{10,11} All of these studies indicate that rHA trimers could be an important component of a subunit vaccine strategy against influenza.

Hemagglutinin (HA)-specific serum antibody titers correlate with protection from disease; however, recombinant proteins are often weak immunogens and require multiple, high-dose immunizations to achieve protection.^{6,7,12} Polyanhydride nanoparticles have been shown to be a versatile vaccine adjuvant/delivery platform capable of enhancing the immune response to recombinant proteins.^{13–15} Polyanhydrides are biodegradable materials that provide sustained release kinetics of encapsulated antigens, resulting in long-lived, high-avidity antibody titers, even with suboptimal doses of antigen.^{13–15} In addition to amplifying humoral immunity, polyanhydride nanoparticle-based vaccines (ie, nanovaccines) have been shown to be immunomodulatory¹⁶ and are capable of promoting cell-mediated immunity. Previous work with polyanhydride nanovaccines indicated expansion of antigen-specific CD8⁺ T cells following nanovaccine immunization, resulting in memory T cell populations that responded to antigen-expressing tumor challenge (CD8⁺ T cells).¹⁷ This broad repertoire of immune responses induced by polyanhydride nanovaccines may prove beneficial for influenza vaccine efficacy because robust cell-mediated responses are often associated with broader protective immunity and directed at conserved epitopes.^{18,19}

Previously, soluble H5 HA trimers (sH5₃) from H5N1 influenza virus A/Whooper Swan/Mongolia/244/05 were produced using a baculovirus insect cell expression system and shown to maintain their oligomeric structure and antigenicity upon release from polyanhydride nanoparticles.²⁰ In this study, the immunogenicity and virus-neutralizing antibody titer in response to nanovaccine immunization were

observed for approximately 2 months, and concluded with an analysis of memory T cell responses. Finally, the efficacy of the nanovaccine formulations was examined using a low-pathogenic, live-viral challenge. The data demonstrate that polyanhydride nanovaccines containing immunogenic HA trimer represent a potentially viable platform for pandemic influenza vaccines.

Materials and methods

Plasmids and antibody

Plasmid pHW500 (Genbank DQ659326), obtained from Dr Bruce Janke of Iowa State University, contains the full-length HA gene from HPAI H5N1 influenza virus A/Whooper Swan/Mongolia/244/05 (H5N1). The pHW500 HA gene was modified by replacement of the cognate polybasic cleavage site with that from a low-pathogenic H6N1 avian influenza virus.²¹ The FDA-VN plasmid, obtained from Dr Carol Weiss of the US Food and Drug Administration, contains the full-length HA from A/Vietnam/1203/2004 (H5N1) (Genbank EF541403) in pCMV/R.²² The pWS-HA was constructed by replacing the H5 Vietnam HA gene in FDA-VN with the HA gene from pHW500. The low-pathogenic cleavage site in pWS-HA was replaced with the polybasic cleavage site from A/Whooper Swan/Mongolia/3/05 (H5N1) (Genbank AB233320.1). The equine infectious anemia lentivirus vector plasmids pEV53B and pSIN6.1ClucW were obtained from John Olsen.²³ Hyperimmune swine sera containing high titers of H5 neutralizing antibody was obtained from Dr Bruce Janke.²¹ H5-specific polyclonal rabbit antisera was obtained from Dr Carol Weiss.²²

Cloning and expression of soluble H5 trimer (sH5₃)

The HA ectodomain (nucleotides 1–1,723) was amplified from pHW500 and modified at the 3' end by addition of linker sequences, a GCN4pII trimerization domain,⁸ and His-tag sequences (Figure 1A). The modified gene was cloned into pFastBac I, transformed into DH10Bac cells, and recombinant baculoviruses were generated using Bacto-Bac Baculovirus Expression System (Thermo Fisher Scientific, Waltham, MA, USA). Sf9 cells (Thermo Fisher Scientific) were infected with recombinant baculovirus and supernatants collected 96 hours after infection were clarified by centrifugation, dialyzed against 10 mM Tris buffer (10 mM Tris, 50 mM NaCl, pH 8.0), and incubated with Ni-NTA beads (Thomas Scientific, Swedesboro, NJ, USA) overnight at 4°C. The beads were washed with 10 mM Tris buffer containing 10 mM imidazole and recombinant

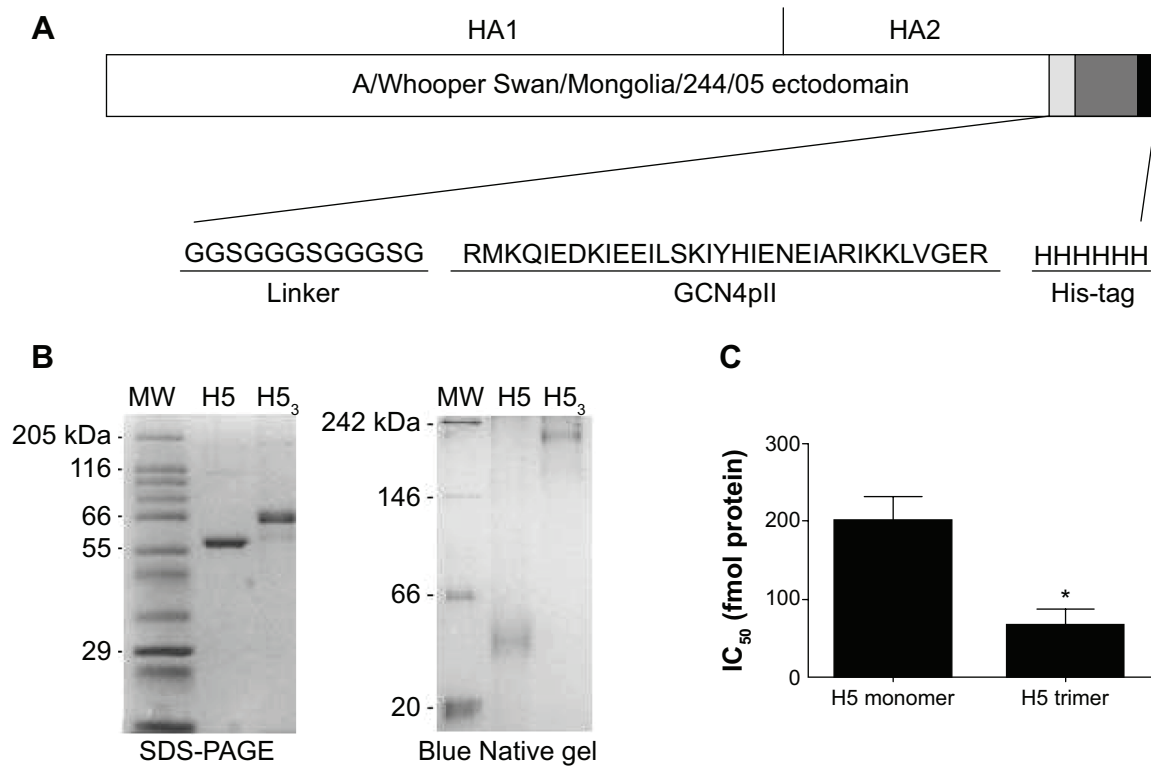


Figure 1 Expression and characterization of H5₃.

Notes: (A) The HA ectodomain (nucleotides 1–1,723) was amplified from pHW500 and modified by addition of linker sequences, a GCN4pII trimerization domain, and His-tag sequences at the 3' end. (B) SDS-PAGE and Blue Native gel electrophoresis confirmed the expression of the H5 monomer and trimeric forms. (C) A neutralization inhibition assay demonstrated that the immunogenicity of trimeric H5 was significantly greater than that of monomeric forms, requiring a lower concentration of protein to inhibit 50% of the neutralizing sera (IC₅₀). Error bars represent the standard error of the mean. *Indicates statistical significance. $P \leq 0.0243$.

Abbreviations: HA, hemagglutinin; H5, H5 hemagglutinin monomer; H5₃, H5 hemagglutinin trimer; SDS-PAGE, sodium dodecyl sulfate polyacrylamide gel electrophoresis; IC₅₀, inhibitory concentration 50%.

sH5₃ was eluted in 10 mM Tris buffer containing 250 mM imidazole. The eluted proteins were dialyzed against 10 mM Tris buffer to remove imidazole, concentrated, and stored at 4°C for up to 1 month or at –80°C for long-term storage.

HA-pseudotyped reporter virus

HA-pseudotyped viruses were produced in HEK293T cells as described previously.^{22,24} Briefly, HEK293T cells were cotransfected with 5 µg of pEV-53B encoding the lentiviral core proteins of equine infectious anemia virus, 5.5 µg of the luciferase reporter plasmid pIgSIN6.1Luc, and 1 µg of pFDA-VN or pWS-HA plasmid DNA expressing HA from A/Vietnam/1203/2004 (clade 1) or A/Whooper Swan/Mongolia/244/05 (clade 2.2), respectively. At 18 hours post-transfection, cells were incubated with fresh medium containing 7 mU/mL of *Vibrio cholerae* Type II neuraminidase (NA) (Sigma-Aldrich Co., St Louis, MO, USA) and 10 mM NaB to induce the release of H5-pseudovirions from the surface of the producer cells. Supernatants were collected 48 hours post-transfection, clarified by centrifugation, and stored at –80°C.

Reporter virus pseudotyped with HA from A/Vietnam/1203/2004 (H5-VN-Luc) or A/Whooper Swan/Mongolia/244/05 (H5-WS-Luc) were titrated on HEK293T cells. Cells were seeded in 96-well plates at 2×10^4 cells/well in Dulbecco's modified Eagle medium supplemented with antibiotics and heat-inactivated fetal bovine serum. The following day, cells were inoculated in triplicate with 10-fold serial dilutions of pseudovirus stock in the presence of 8 µg/mL polybrene. At 48 hours post-transduction, cells were lysed and assayed for luciferase activity using the ONE-Glo Luciferase assay system (Promega Corporation, Fitchburg, WI, USA). Luciferase activity was quantified using a Centro XS3 LB960 illuminometer (Berthold Technologies, Bad Wildbad, Germany) and results reported as relative light units (RLU)/mL supernatant.

Electrophoresis

For protein analyses in denaturing conditions, 1 µg of purified sH5₃ protein was boiled for 5 minutes in sodium dodecyl sulfate (SDS) loading buffer (50 mM Tris, 1% β-mercaptoethanol, 2% SDS, 0.005% bromophenol blue,

and 10% glycerol) and electrophoresed in 10% SDS-PAGE (polyacrylamide gel electrophoresis). For analyses in non-denaturing gels, 3 µg of purified sH5₃ protein was mixed with Blue Native loading buffer (2 mM EDTA, 20 mM NaCl, 20 mM Bis-Tris, 10% glycerol, and 0.08% Coomassie Blue G-250) and separated on 10% Blue Native PAGE gel containing Bis-Tris, glycerol, and acrylamide in Bis-Tris buffer in the outer chamber and Tricine, Bis-Tris with Coomassie Blue G250 in the inner chamber. Following electrophoresis, gels were stained with Coomassie Blue and imaged with a GelDoc XR+ imaging system (Bio-Rad Laboratories Inc., Hercules, CA, USA).

Polymer and nanoparticle synthesis

Diacids based on 1,8-bis(*p*-carboxyphenoxy)-3,6-dioxaoctane (CPTEG) and 1,6-bis(*p*-carboxyphenoxy)hexane (CPH) were synthesized as previously described^{25,26} using 1,6-dibromohexane, 1-methyl-2-pyrrolidinone, hydroxybenzoic acid, N,N-dimethylacetamide, and tri-ethylene glycol (Sigma-Aldrich Co.); acetic acid, acetone, acetonitrile, dimethyl formamide, potassium carbonate, sulfuric acid, and toluene (Thermo Fisher Scientific, Waltham, MA, USA); 4-*p*-fluorobenzonitrile (Apollo Scientific, Bredbury, UK). Following diacid synthesis, 20:80 CPTEG:CPH copolymer was synthesized by melt polycondensation.²⁶ The final composition of the polymer (23:77 CPTEG:CPH), molecular weight (6,055 Da), and purity were determined with ¹H nuclear magnetic resonance spectroscopy (VXR 300 MHz; Varian, Palo Alto, CA, USA). H5₃-loaded nanoparticles were synthesized using a water-oil-oil double emulsion process.²⁷ First, 4.5 mg of sH5₃ was concentrated into 100 µL of nanopure water. The solution of sH5₃ was then added to 45 mg of 20:80 CPTEG:CPH dissolved in 2.25 mL of methylene chloride and homogenized for 90 seconds. The H5₃-loaded nanoparticles were precipitated by pouring the solution into 562.5 mL of chilled pentane and collected via vacuum filtration. Blank nanoparticles and 1% polyinosinic-polycytidylic acid (poly

I:C) (InvivoGen, San Diego, CA, USA)-loaded nanoparticles were synthesized similarly without water. Scanning electron microscopy (Quanta 250; FEI, Hillsboro, OR, USA) was used to characterize the size and morphology of the nanoparticles, which were found to be consistent with previous work.¹³ The encapsulation efficiency of protein was determined by degrading 10 mg of nanoparticles in 250 µL of 40 mM sodium hydroxide and quantifying the total protein released using a microBCA protein kit (Pierce, Rockford, IL, USA).¹³

Mice

Female BALB/c mice were purchased from Harlan Laboratories (Indianapolis, IN, USA). All mice were housed under specific pathogen-free conditions where all bedding, caging, water, and feed were sterilized prior to use. Animal procedures were conducted with the approval of the Iowa State University and University of Nebraska Medical Center Institutional Animal Care and Use Committees.

Immunizations

For each formulation, mice were administered a total of 10 µg H5₃. As outlined in Table 1, a portion of the 10 µg H5₃ dose was encapsulated into 50 µg polyanhydride nanoparticles, and delivered with or without 250 µg nanoparticles encapsulating poly I:C. Blank (ie, no encapsulated protein or poly I:C) nanoparticles were used to keep the total amount of nanoparticles (300 µg) constant among the different vaccine regimens. In addition, control formulations composed of 10 µg of soluble H5₃ (sH5₃) alone, 10 µg of sH5₃ delivered with blank nanoparticles, or 10 µg of sH5₃ in conjunction with 10 µg of monophosphoryl lipid A (MPLA) from *Salmonella enterica* (Sigma-Aldrich Co.) were administered. All formulations were suspended in 250 µL (subcutaneous [SC] immunization) or 50 µL (intranasal [IN] immunization) of sterile saline. Formulations containing nanoparticles were sonicated for 30 seconds to ensure dispersion of particle aggregates before immunization. SC immunizations were administered

Table 1 H5₃ vaccine formulations

Group	Encapsulated H5 ₃	Soluble H5 ₃	Total 20:80 CPTEG:CPH nanoparticles	Additional excipients
H5 ₃ -nanoparticles	2 µg	8 µg	300 µg	–
H5 ₃ - & poly I:C-nanoparticles	2 µg	8 µg	300 µg	2.5 µg poly I:C
Blank nanoparticles	–	10 µg	300 µg	–
MPLA	–	10 µg	–	10 µg MPLA
sH5 ₃	–	10 µg	–	–
Saline	–	–	–	–

Abbreviations: CPH, 1,6-bis(*p*-carboxyphenoxy)hexane; CPTEG, 1,8-bis(*p*-carboxyphenoxy)-3,6-dioxaoctane; H5₃, H5 hemagglutinin trimer; MPLA, monophosphoryl lipid A; sH5₃, soluble H5 hemagglutinin trimer.

at the nape of the neck; IN immunizations were carried out using droplet admission via pipettor after administration of ketamine/xylazine chemical anesthetic. For “prime/boost/boost” regimens, booster immunizations were prepared and administered the same way as primary immunizations at days 21 and 42. Serum samples were obtained at the time points indicated via saphenous vein bleeding.

Neutralization assay

Neutralizing antibody assays were carried out using HA-pseudotyped reporter virus as described previously.^{22,24} HEK293T cells were seeded in a 96-well plate at 2×10^4 cells/well and grown for 24 hours. Sera samples were serially diluted three-fold in culture medium containing 8 $\mu\text{g/mL}$ polybrene (Sigma-Aldrich Co.), and mixed with an equal volume of diluted pseudovirus stock containing 5×10^4 RLU/mL. After incubation at 37°C for 1 hour, virus and serum mixtures were added to the cells. Infectivity was evaluated 48 hours after transduction using the One-glo Luciferase assay system (Promega Corporation). The percentage neutralization was calculated as $(1 - [\text{virus} + \text{sera RLU}/\text{virus only RLU}]) \times 100$. The percentage neutralization for each sera dilution was plotted and neutralizing titers were reported as infectious dose 50 (ID_{50}), calculated as the reciprocal of the serum dilution that neutralized 50% of the virus.

A neutralization inhibition assay was used to evaluate the immunogenic properties of sH5₃ protein. Ten-fold serial dilutions of sH5 trimer or monomer were incubated at 37°C for 30 minutes with a 1:5,000 dilution of pig H5-antiserum (Dr Bruce Janke, Iowa State University). Following pre-incubation with H5 protein, 50,000 RLU of H5-WS-Luc was added and the samples incubated for an additional 1 hour at 37°C and inoculated onto HEK293T cells in 96-well plates. At 48 hours after transduction, the levels of luciferase in the transfected cells were evaluated using the One-Glo luciferase assay system (Promega Corporation) and percentage neutralization was calculated as described above. The percentage neutralization at each protein concentration was plotted and the half maximal inhibitory concentration (IC_{50}) was calculated as the concentration of soluble H5 protein that inhibited 50% of the neutralizing activity of the diluted pig sera.

Flow cytometry for T cell memory populations

Brachial and axillary lymph nodes were harvested 63 days postimmunization and homogenized into single cell suspensions in complete tissue culture medium. Single cell populations were labeled with 2.5 μM 5-(and-6)-carboxyfluorescein

diacetate, succinimidyl ester (5[6]-CFDA, SE; CFSE; Thermo Fisher Scientific, Waltham, MA, USA). Cells (2.5×10^5) were incubated in 96-well U-bottom plates with 0.5 μg of sH5₃ antigen for 96 hours. Cells were aspirated and cell surface marker expression and proliferation were evaluated using flow cytometry (BD FACSCanto; BD Biosciences, San Jose, CA, USA). Cell suspensions were blocked for nonspecific antibody binding using 0.1 mg/mL Rat IgG (Sigma-Aldrich Co.) and 10 $\mu\text{g/mL}$ mouse anti-CD16/32 (eBioscience, San Diego, CA, USA). Fluorescently conjugated antibodies specific for murine CD4 and CD8 (eBioscience) were diluted in FACS buffer and were used to quantify specific cell populations.

Low-pathogenic viral challenge and clinical evaluation

The efficacy of H5₃-based nanoparticle vaccines were evaluated in mice that were challenged at 63 days postimmunization with the low pathogenicity influenza virus A/H5N1 VN/H5N1-PR8CDC-RG, obtained from the Centers for Disease Control and Prevention (Atlanta, GA, USA). This is a PR-8-derived reassortant virus that contains the HA and NA genes of H5N1 A/Vietnam/1203/04, a clade 1 virus. Mice were anesthetized with 20 mg/mL of xylazine and 100 mg/mL of ketamine (1:4 ratio) and inoculated intranasally with 2.7×10^3 tissue culture infectious doses 50% (TCID_{50}) virus in 30 μL phosphate-buffered saline. A control group consisting of naïve (ie, no vaccine), no challenge (ie, no infection) mice were housed similarly throughout the duration of the study. Three days after challenge, half of the mice were euthanized with 600 μL of 20 mg/mL xylazine and 100 mg/mL ketamine delivered via intraperitoneal injection. Bronchoalveolar lavage (BAL) fluid was collected as described previously²⁸ and lung tissue was collected for inflammatory cytokine and viral load quantitation (described below). The remaining mice were monitored for weight loss for 2 weeks postchallenge before being removed from study.

Virus load quantitation

Following the procedure of Alsharifi et al lung tissue was preserved in 3 mL of RNeasy Lysis Reagent (Qiagen NV, Venlo, the Netherlands).²⁹ Tissue was held submerged in the RNeasy Lysis Reagent for 3 days at 4°C . Tissues were then removed from the RNeasy Lysis Reagent, weighed, and cut into 30 mg pieces and individually frozen at -80°C in 1.5 mL microcentrifuge tubes. For the extraction process (total RNA), individual 30 mg tissue pieces were homogenized in 200 μL buffer RLT (RNeasy Mini Kit, Qiagen NV) using a disposable pellet pestle (Thermo Fisher Scientific) in conjunction with

a cordless motor (Thermo Fisher Scientific). An additional 400 μL of buffer RLT was added to each tube after homogenization was completed. Tissue was extracted into 60 μL final volume in sterile, RNase-free H_2O (Qiagen NV) and frozen at -80°C until polymerase chain reaction (PCR) was performed. Samples containing the extracted RNA were thawed, mixed well, and the total RNA concentration was determined, in duplicate measurements, using the Nanodrop method for RNA content. Total RNA concentration for each sample was adjusted to 40 $\mu\text{g}/\mu\text{L}$, and 5 μL was used as the template for the PCR reaction. PCR was performed on an Applied Biosystems 7500 Fast Real-Time PCR System, on the standard mode, using AgPath-ID One-Step RT-PCR Reagents (Thermo Fisher Scientific, Waltham, MA, USA) in conjunction with the Fast-Track Diagnostics FTD-21-96/12 Kit (Junglinster, Luxembourg), which contains bome mosaic virus (BMV) internal PCR extraction control, positive control, and primer/probes for universal influenza A antigen. For the standard curve, normal, non-influenza-challenged mice lungs (naïve controls) were homogenized using the procedure outlined above. RNA from stock influenza A H5 virus was extracted using the Qiagen QiAmp Viral RNA Kit Mini Kit (Qiagen NV). Extracted RNA was quantified using the Nanodrop procedure. For the standard curve, ten-fold dilutions of the H5-extracted viral RNA were mixed with extracted RNA from the normal mouse lungs that had been standardized to 40 $\text{ng}/\mu\text{L}$. Standard curves were obtained with each set of PCR reactions.

Cytokine analysis of BAL fluid

BAL fluid samples collected 3 days post-viral challenge were analyzed for the inflammatory cytokines IL-6, IP-10, MIG, G-CSF, IFN- γ , MCP-1, KC, and MIP-2 using a MILLIPLEX[®] MAP assay kit (EMD Millipore, Billerica, MA, USA). The assay was performed according to manufacturer instructions. Briefly, 25 μL of BAL fluid, 25 μL assay buffer, and 25 μL MILLIPLEX MAP beads were added to each well of a 96-well plate. After shaking overnight at 4°C , the plate was washed and incubated with 25 μL /well detection antibody for 1 hour at room temperature. Following, 25 μL /well of streptavidin phycoerythrin was added for an additional 30 minutes. The plate was washed once more before measuring fluorescence intensity on a Bio-Plex 200 system (Bio-Rad Laboratories Inc.).

Principal component analysis

Previously, principal component analysis (PCA) was used to identify pathogen-mimicking capabilities of polyanhydride nanoparticles in comparison to bacteria.³⁰ In this study, PCA

was used to mathematically capture the similarities among the different vaccine formulations tested and their immune responses. A linear data dimensionality reduction method^{31–33} was used to condense eight-dimensional data (neutralizing antibody titer, viral load, and six cytokine concentrations) into two dimensions (ie, principal components [PC]). In this work, PC1 was a linear combination of viral load and cytokine concentration(s), while PC2 primarily captured neutralizing antibody titer. At least 90% of the variance was captured between these two PCs. Similarities among data were mathematically represented as the proximity between data points (ie, data points close together are more similar than data points farther apart).

Statistics

Statistical significance among formulations ($P \leq 0.05$) was determined by one-way analysis of variance (ANOVA) followed by Tukey's post-test using Graph Pad Prism (v6.01; Graph Pad Software, Inc., La Jolla, CA, USA).

Results

Characterization of soluble H5 trimer protein (sH5₃)

A recombinant baculovirus containing the ectodomain of HA from A/Whooper Swan/Mongolia/244/05 (Figure 1A) was expressed in Sf9 insect cells and the secreted soluble H5 trimer (sH5₃) protein was purified from culture supernatant by affinity chromatography using nickel agarose beads. Analyses of the purified protein by electrophoresis in SDS-PAGE and Blue Native gels indicated the predominant form of sH5₃ protein was trimeric (Figure 1B). A neutralization inhibition assay was used to determine whether the sH5₃ retained neutralizing epitopes of native virus (Figure 1C). In this assay, convalescent pig sera containing high titers of H5-neutralizing antibody was incubated with serial dilutions of sH5₃ or sH5 monomer and tested for neutralizing activity against H5-WS-Luc pseudotyped reporter virus. The HA presented in its native trimeric state has been found to be more immunogenic than the monomeric form.^{5,8} Similarly, we observed that while both the monomeric and trimeric forms of sH5 inhibited neutralizing antibody at fmol concentrations, the inhibitory activity of the sH5₃ trimer was about 2.5-fold higher than that of the monomer.

Immunogenicity of soluble recombinant H5 HA trimer

BALB/c mice (6–8 weeks) were immunized either subcutaneously or intranasally with 10 μg of soluble protein at 0, 21, and 42 days and sera collected at 21, 42, and 63 days

postimmunization were tested for neutralizing antibody against the homologous H5-WS-Luc pseudovirus. No neutralizing antibody was detected at day 21 following primary immunization (Figure 2A). By day 42, the majority of mice in both immunization groups had detectable neutralizing antibody titers (ie, >50) that were greater than protective titers reported in the literature.^{7,8} However, the response was highly variable among individual mice, with ID_{50} titers ranging from undetectable to greater than 1,000. Neutralization titers significantly increased following a second boost (day 63), with mean neutralization antibody titers of greater than 10,000. In addition, there was less variability in the neutralizing antibody titers among individual mice, especially those immunized subcutaneously.

As highly pathogenic H5N1 has continued to spread in avian and mammalian hosts, different lineages have emerged that are now classified into distinct, but phylogenetically related clades.¹ To examine the cross-clade breadth of the neutralizing antibody response elicited with A/Whooper Swan/Mongolia/244/05-based sH5₃ (clade 2.2), day 63 sera was tested for neutralizing activity against H5-VN-Luc, a luciferase reporter virus pseudotyped with HA from the A/Vietnam/1203/2004 HA, an H5 clade 1 strain. Although the cross-clade neutralization titers were 2–10-fold lower than titers to homologous virus, serum samples from all mice were able to neutralize Vietnam/1203 pseudotyped virus, with mean neutralizing antibody titers of 1,000 (Figure 2B). Together, these data indicate that nonadjuvanted sH5₃ antigen is able to elicit high titers of H5 cross-clade neutralizing antibody when administered using a prime/boost/boost regimen. Consistent with previous reports,^{6,7,12} multiple immunizations

of sH5₃ recombinant protein were required to generate consistently high neutralizing antibody titers.

Neutralizing antibody response in mice vaccinated with H5₃-based nanovaccines

Our previous studies showed that polyanhydride nanovaccines enhanced the immune response to recombinant proteins.^{13–15} Therefore, we examined the immune response of sH5₃ delivered with and/or encapsulated into polyanhydride nanoparticles (Table 1). Mice were immunized subcutaneously with either a single-dose regimen or a prime/boost/boost regimen of three immunizations 21 days apart. Serum was collected at 42 and 63 days postimmunization and tested for neutralizing antibody titers against H5-WS-Luc. As we observed above, sH5₃ alone elicited low neutralizing titers in the absence of booster immunizations, with most mice having ID_{50} titers of less than 100 through 63 days postimmunization (Figure 3). Higher titers were observed using a single-dose immunization regimen with sH5₃ in the presence of adjuvant, with MPLA eliciting higher neutralizing antibody titers than any of the nanovaccine formulations or sH5₃ alone. In the absence of booster immunizations, the neutralizing antibody titers in sera from mice immunized with nanovaccine formulations was lower than that from the MPLA-treated mice at 42 days, with the exception of the formulation including poly I:C; however, there were a greater number of single-dose, nanoparticle-vaccinated mice with detectable neutralizing antibodies than in mice vaccinated once with sH5₃ alone. Administration of one or two booster vaccinations resulted in enhanced neutralizing antibody titers in all the formulations. The highest and most consistent titers were observed after

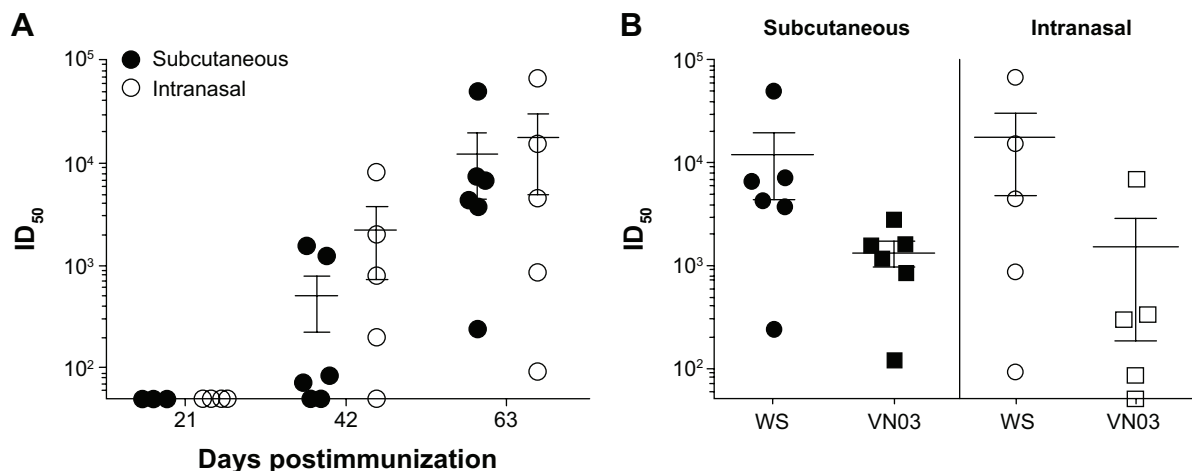


Figure 2 Immunogenicity of soluble H5₃ antigen.

Notes: (A) Mice were immunized subcutaneously or intranasally with 10 μ g soluble H5₃ on days 0, 21, and 42. Sera collected at days 21, 42, and 63 after primary immunization were tested for neutralizing antibody. Neutralizing antibody titers were not detected until after the second immunization; however, titer generated by days 42 and 63 were robust. (B) The neutralizing antibody titers demonstrated cross-clade protection at 63 days with both immunization routes. Error bars represent the standard error of the mean.

Abbreviations: H5₃, H5 hemagglutinin trimer; ID_{50} , sera dilution that inhibits 50% of pseudovirus infectivity.

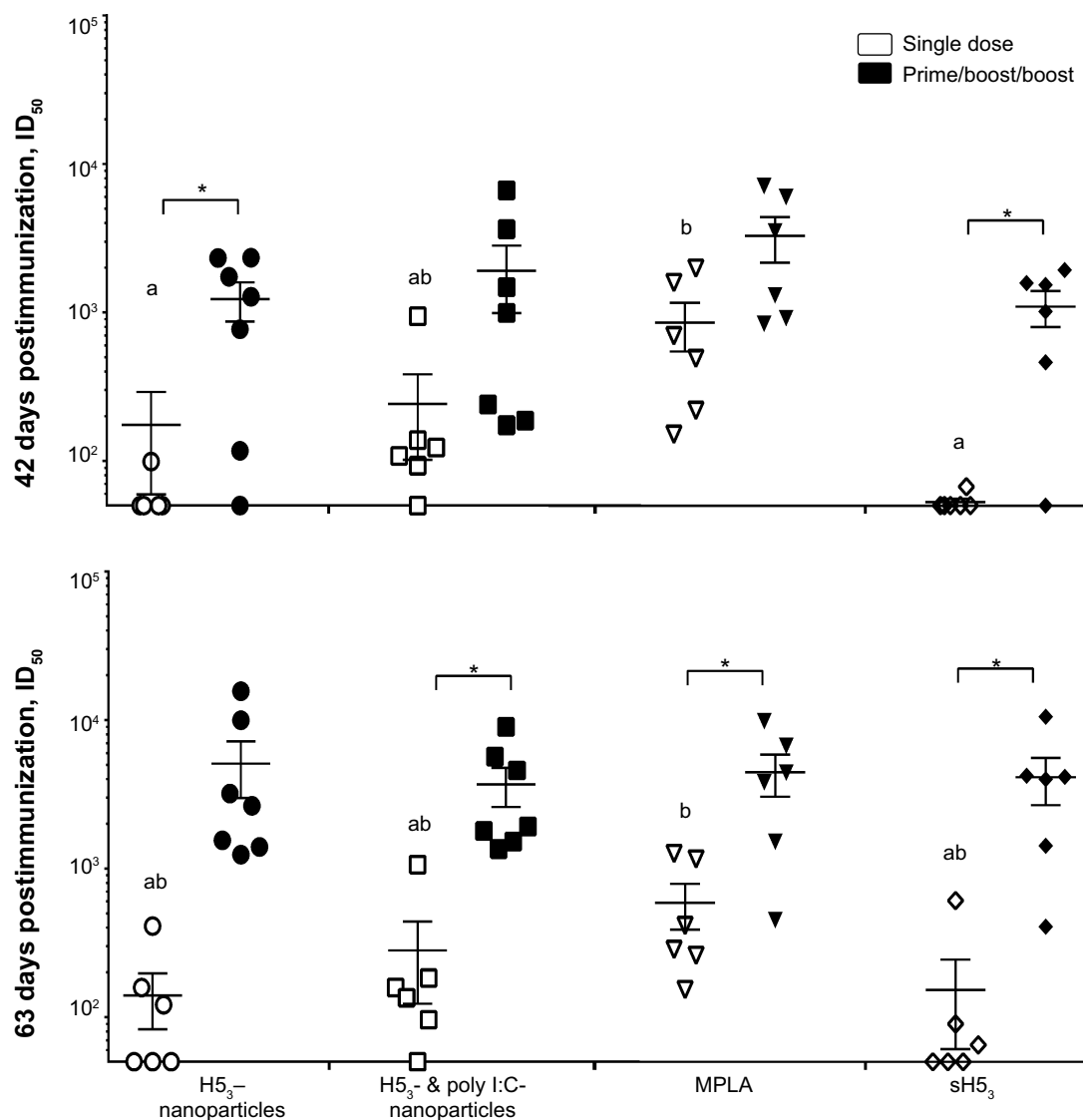


Figure 3 The neutralizing antibody responses to H₅₃-polyanhydride nanovaccines were examined 42 and 63 days post-primary immunization.

Notes: Single-dose vaccination regimens utilizing poly I:C encapsulated with H₅₃ demonstrated neutralization titers similar to a MPLA-adjuvanted control at day 42, with most of the remaining formulations reaching equivalency at day 63. The use of a prime/boost/boost immunization regimen enhanced the production of neutralizing antibodies and reduced mouse-to-mouse variability. Error bars represent the standard error of the mean. Different letters indicate statistical significance among treatments of the same regimen (ie, single dose or prime/boost/boost) at each time point. *Indicates statistical significance between the different regimens for each treatment. No statistical significance was observed when comparing between the various prime/boost/boost formulations. $P \leq 0.0434$.

Abbreviations: H₅₃, H5 hemagglutinin trimer; MPLA, monophosphoryl lipid A; sH₅₃, soluble H5 hemagglutinin trimer; ID₅₀, sera dilution that inhibits 50% of pseudovirus infectivity; poly I:C, polyinosinic-polycytidylic acid.

receiving three immunizations (prime/boost/boost regimen, day 63), with little difference observed among the different formulations (Figure 3).

Cell-mediated immune response

Cell-mediated immune responses are often associated with broader protective immunity and can play an important role in protection against antigenically diverse strains of influenza.^{18,19} Therefore, we examined the proliferative T cell populations on day 63 (ie, 21 days after the last immunization) in mice receiving the prime/boost/boost regimen (Figure 4). Draining lymph nodes were removed and homogenized to single cell

suspensions and labeled with CFSE to observe proliferating cell populations. After 96 hours of ex vivo stimulation with sH₅₃, CD4⁺ T cells were quantified via flow cytometry. A significant expansion of CD4⁺ T cells was observed in all groups of vaccinated mice as compared to mice receiving saline alone. In addition, the inclusion of poly I:C within the nanovaccine formulation resulted in significantly higher numbers of CD4⁺ T cells as compared to mice immunized with sH₅₃ alone or sH₅₃ adjuvanted with MPLA. These data suggest that inclusion of nanoparticles encapsulating poly I:C to induce an appropriate innate immune response may enhance antigen-specific adaptive immune responses.

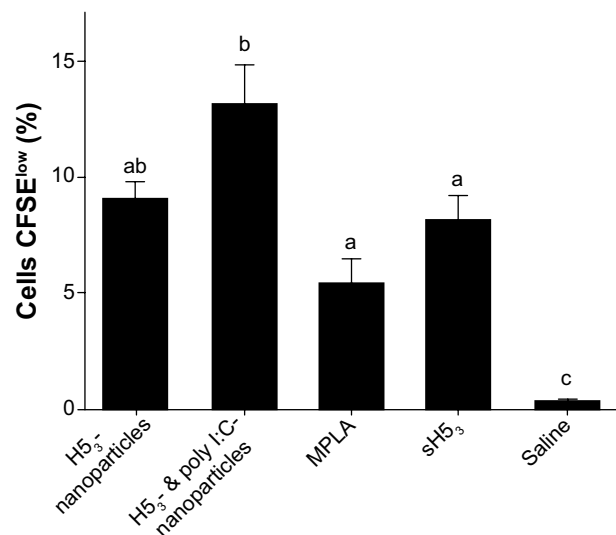


Figure 4 Enhanced CD4⁺ T cell memory with poly I:C polyanhydride nanovaccine. **Notes:** Draining lymph nodes were harvested from prime/boost/boost immunized mice 63 days after the primary immunization. Ex vivo antigen stimulation and CFSE-labeling demonstrated enhanced CD4⁺ T cell proliferation of mice immunized with poly I:C nanovaccines. Error bars represent the standard error of the mean. Different letters indicate statistical significance among treatments. $P \leq 0.0147$.

Abbreviations: CFSE, 5-(and-6)-carboxyfluorescein diacetate, succinimidyl ester; MPLA, monophosphoryl lipid A; sH5₃, soluble H5 hemagglutinin trimer; poly I:C, polyinosinic-polycytidylic acid.

H5₃ vaccination protects against live-viral challenge

To examine the efficacy of H5₃-loaded nanoparticles in protection from clinical disease, mice were vaccinated with various nanovaccine formulations (Table 1) using a prime/boost/boost regimen and challenged at day 63 with the low-pathogenic VNH5N1-PR8/CDC-RG, a reverse genetics-derived influenza virus containing the HA and NA genes of A/Vietnam/1203/04 (H5N1) virus in the genetic background of the high-growth master strain A/Puerto Rico/8/34 (H1N1). Pre-challenge antibody response to the various vaccine regimens was evaluated in serum collected 1 week before challenge. All vaccinated groups had high levels of total anti-H5₃ IgG, with no significant difference among the groups. Similar to our earlier studies, all vaccinated mice had detectable neutralizing antibody titers of greater than 100 to the clade 2.2 H5-WS-Luc pseudotyped reporter virus, with mean titers among the group ranging between 3,000–4,000 (Figure S1).

The body weight of each mouse was observed for 14 days postchallenge. All of the vaccine formulations protected mice from challenge with body weight increasing postchallenge similar to naïve mice that were not challenged (Figure 5A). In contrast, mice receiving saline immunizations began to lose a significant proportion of their body weight approximately 5 days postchallenge. Saline-administered

mice continued to lose approximately 20% of their total weight by 8 days postchallenge before recovering. Based on clinical response, as measured by weight loss, all vaccine regimens induced a protective immune response and there was no significant difference among the various vaccine regimens.

At 3 days postchallenge, half of the mice in each group were sacrificed and lung homogenates were assayed for virus load using quantitative reverse transcription PCR as described previously.³⁴ All vaccinated mice showed significant reduction in virus load upon challenge as compared to the virus load in the saline-administered mice (Figure 5B). A wide mouse-to-mouse variability in virus load was observed for all vaccine regimens, with no significant differences among the groups in mean virus loads. However, a correlation between pre-challenge (day 56 post-primary immunization) neutralizing antibody titer and virus load (day 3 postchallenge) was observed (Figure 5C).

Finally, a multiplexed assay was used to quantitate the concentrations of inflammatory cytokines (IL-6, IP-10, G-CSF, IFN- γ , MCP-1, KC, and MIP-2) present in BAL fluid. Vaccinated mice had low levels of inflammatory cytokines 3 days postchallenge similar to the naïve, no challenge controls (Figure S2). In contrast, mice receiving saline had significantly greater concentrations of most inflammatory cytokines postchallenge with the exception of IFN- γ (data not shown).

Clinical signs postchallenge in mice vaccinated with H5₃-nanoparticles are similar to healthy naïve, noninfected mice

PCA was utilized to capture the complex relationships between the neutralizing antibody titer, viral load, and six inflammatory cytokine measurements discussed above. Following infection, saline-administered mice (ie, immunologically naïve) had the most unique behavior because this was the only treatment group with any significant PC1 values (Figure 6A). The PC1 dimension captured the largest amount of independent data and was a linear combination of viral load and cytokine concentration. In contrast, the PC2 dimension was primarily captured by neutralizing antibody titer. While all of the H5₃ vaccine formulations demonstrated similar PC1 values, mice receiving the 'H5₃- & poly I:C-nanoparticle' formulation or 'sH5₃' showed the most significant PC2 values, or variability in neutralizing antibody among animals. Therefore, saline-administered mice have the behavior most unlike the remaining formulations, and neutralizing antibody titer is the most unique of the descriptors used in the analysis.

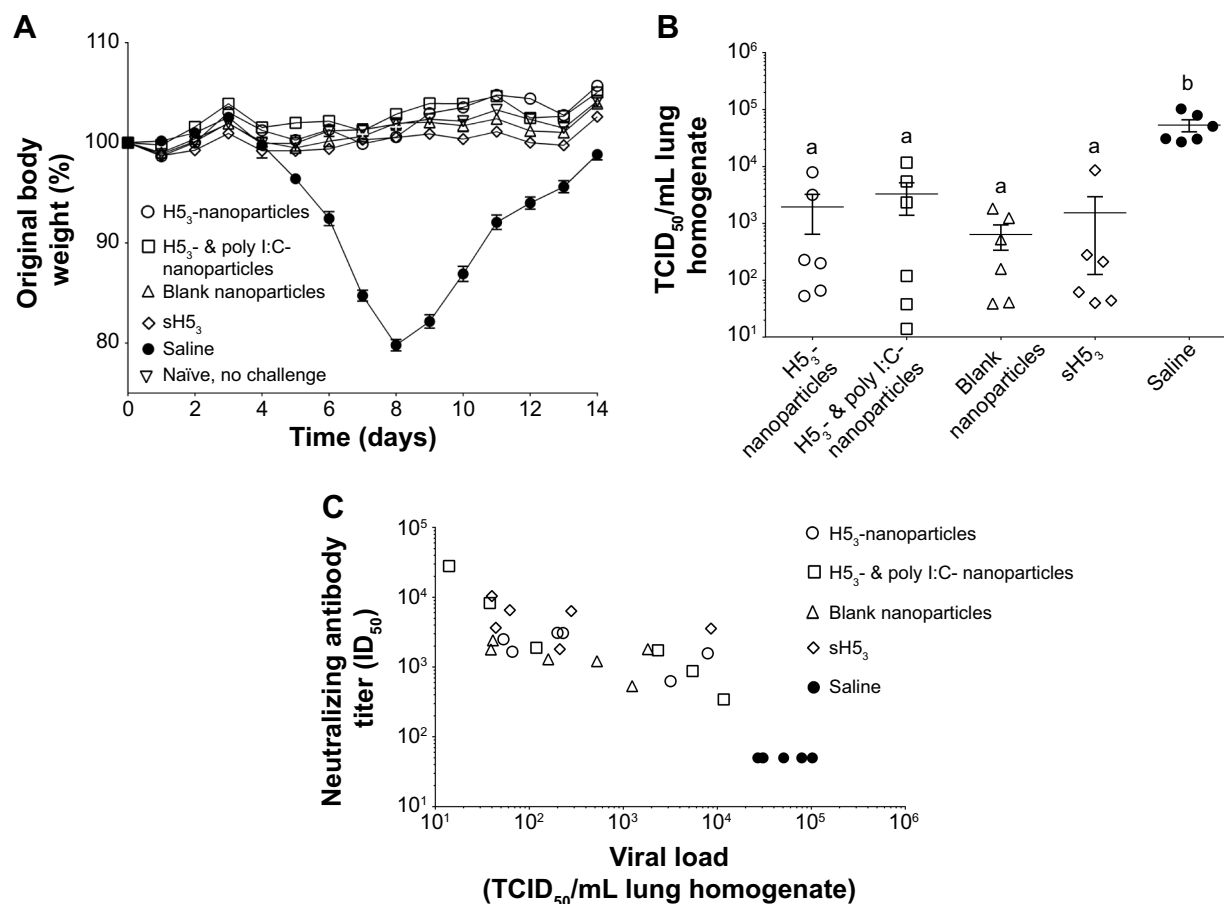


Figure 5 Mice were protected upon challenge with low pathogenic VN5N1-PR8/CDC-RG.

Notes: (A) Mice were challenged with a reverse genetics-derived influenza virus, VN5N1-PR8/CDC-RG, containing the HA and NA genes of A/Vietnam/1203/04 (H5N1) virus in the genetic background of the high-growth master strain A/Puerto Rico/8/34 (H1N1). Body weight was observed for 2 weeks postinfection. All vaccinated mice maintained or gained weight similar to healthy naïve, no challenge mice. (B) Viral load in lung homogenates collected at 3 days post-challenge, (C) which correlated with the pre-challenge neutralizing antibody titers. Different letters indicate statistical significant among treatments.

Abbreviations: H5₃, H5 hemagglutinin trimer; HA, hemagglutinin; ID₅₀, sera dilution that inhibits 50% of pseudovirus infectivity; NA, neuraminidase; sH5₃, soluble H5 hemagglutinin trimer; TCID₅₀, tissue culture infectious dose 50%.

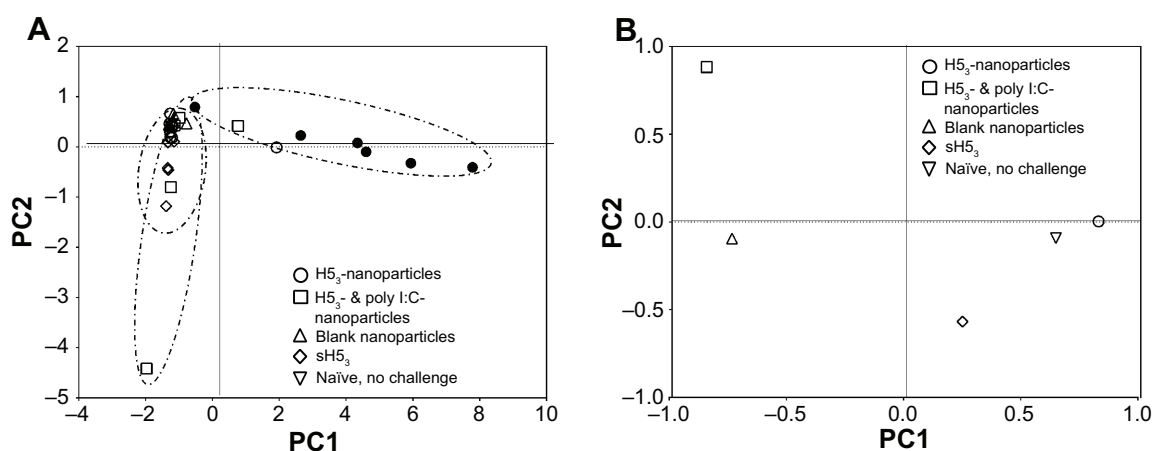


Figure 6 H5₃-nanoparticle responses most similar to healthy naïve, no challenge mice.

Notes: (A) Principal component analysis was performed utilizing neutralizing antibody titer, viral load, and inflammatory cytokine measurements, with each data point representing one mouse. All vaccination formulations were found to be very different from saline-administered mice, as evidenced by the small and high PC1 values, respectively. (B) PCA displaying average values after the exclusion of saline-administered mice and neutralizing antibody titers. The H5₃-nanoparticle formulation induced responses very similar to healthy, noninfected mice.

Abbreviations: H5₃, H5 hemagglutinin trimer; PC, principal components; PCA, principal component analysis.

In addition to the primary correlations uncovered by correlations to saline-administered mice and neutralizing antibody titers, more subtle relationships in the data remained hidden. Thus, the analysis was repeated after excluding the primary correlations (ie, saline-administered mice and neutralizing antibody titer) and is shown in Figure 6B, where only the average value of each treatment is displayed for clarity. The 'H5₃-nanoparticle' formulation induced a response postchallenge similar to the healthy naïve, nonchallenged (ie, control) mice, as seen by the close proximity of the average data points (Figure 6B). Mice receiving soluble H5₃ alone (sH5₃) also resulted in responses similar to the control mice, but less so than the 'H5₃-nanoparticle' formulation. This analysis demonstrated that all the vaccine formulations resulted in a significant improvement in clinical responses when compared to the saline-treated, infected mice, while the 'H5₃-nanoparticle' formulation induced responses most similar to healthy control mice.

Discussion

The design of effective vaccines for the prevention of influenza is a challenging task. Despite best efforts, the 2013–2014 US seasonal influenza vaccine was shown to be ~60% effective in patients.³⁵ In the case of pandemic influenza, such as H5N1, the efficacy of vaccines can be even further reduced. Currently, the approved H5N1 influenza vaccine consists of two large doses (90 µg) administered 28 days apart, but is only 45% effective.³⁶ In this work, we examined an alternative to conventional influenza vaccines through the use of polyanhydride nanoparticles. By encapsulating H5 antigen into polyanhydride nanoparticles, the foundation has been laid for the design of pandemic influenza vaccines with improved efficacy by demonstrating reduced variability in the neutralizing antibody titer, enhanced T cell responses, and greater stability.²⁰

As mentioned above, the recent seasonal influenza vaccine is approximately 60% effective; however, this estimate has a 95% confidence interval of 5%–68%,³⁵ which is a substantial range that indicates the large (and unacceptable) variability of the host responses to the vaccine. Variability of vaccine effectiveness can be influenced by a large number of factors, including age, obesity, nutrition, or other disease(s) present in the host.^{37–39} While the vaccine formulations used in this work performed similarly, one of the strengths of the polyanhydride nanovaccine is illustrated in its ability to reduced immune response variability. Immunizations with H5₃-nanoparticles were shown to induce more consistent neutralizing antibody titers with less variability compared

to other formulations (Figures 3 and S1). In addition, PCA, which incorporated the neutralizing antibody titer, viral load, and inflammatory cytokine data postchallenge, indicated that H5₃-nanoparticle vaccination led to consistent immune responses (Figure 6A). Similarly, H5₃-nanoparticle immunized mice did not exhibit clinical signs of disease postchallenge and were the most similar to healthy (naïve, noninfected) mice (Figure 6B).

Aside from variability in immune responses, many vaccines struggle to stimulate T cell immunity. Although humoral immunity is often sufficient to provide protection against homologous strains of influenza, cell-mediated immunity is necessary when confronting heterologous variants.¹⁸ CD8⁺ T cells secrete antiviral cytokines and often have a broader epitope spectrum than that of the humoral response.^{18,19} While not fully understood, CD4⁺ T cells play a role in protection against heterologous influenza infection by producing cytokines to restrict viral replication and even providing some lytic activity as well.^{18,19,40} Vaccines that can stimulate both humoral and cell-mediated immunity would likely have an advantage in cross-clade protection. The H5₃ antigen used in this work demonstrated the induction of cross-reactive neutralizing antibodies when delivered alone (Figure 2). However, when delivered in the context of polyanhydride nanoparticles, CD4⁺ T cell proliferation was enhanced upon ex vivo stimulation with antigen (Figure 4). The nanovaccine formulation containing poly I:C induced the greatest proliferation of CD4⁺ T cells. It is known that poly I:C signaling through TLR3 induces the production of several Th1 and Th2 cytokines, as well as type I interferons,⁴¹ which may explain the T cell expansion observed in Figure 4.

Finally, we believe that the ability to encapsulate influenza antigens into polyanhydride nanoparticles will provide additional benefits, including dose sparing and enhanced shelf-life characteristics.^{15,42} While many of the nanoparticle formulations used in this study performed similarly to soluble antigen delivered alone, it is likely that the 10 µg dose of the highly immunogenic H5₃ antigen masked the advantages provided by the polyanhydride nanoparticles. Previously, it has been demonstrated that encapsulation of a suboptimal dose of antigen into polyanhydride particles induced similar antibody titers to administration of 64-fold higher soluble antigen alone.¹⁵ The ability to reduce antigen per dose (and therefore increase vaccine availability) and optimize single-dose formulations will be paramount in providing rapid protection upon a pandemic outbreak of H5N1. Finally, polyanhydride nanovaccines exhibit release of immunogenic protein even when stored at room temperature for 1 year.⁴²

This ability for a vaccine to be stored at room temperature would be a major benefit in terms of maintaining stockpiles of pandemic vaccines.

Conclusion

The studies presented herein demonstrate the strong immunogenic properties of the recombinant H5₃ antigen developed for these studies as well as the immune responses to H5₃-loaded polyanhydride nanovaccines. All formulations were able to induce detectable mean neutralizing antibody titers 42 days postvaccination. The formulation including poly I:C also conferred the additional advantages of enhanced antibody titers at earlier time points and induction of proliferative T cell responses. All vaccine formulations were able to induce protection against a low-pathogenic, live-viral challenge. Of interest, PCA demonstrated that after viral challenge, the mice immunized with the H5₃-nanoparticle formulation were as healthy as naïve, nonchallenged mice and this formulation induced less variable immune responses. The current studies provide a foundation to further exploit the advantages of polyanhydride nanoparticle-based subunit vaccines against influenza.

Acknowledgments

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Disclosure

The authors report no conflicts of interest in this work.

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Supplementary materials

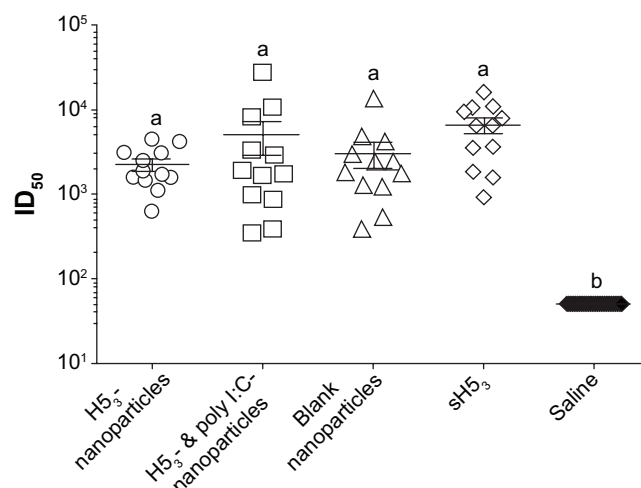


Figure S1 Prime/boost/boost immunized mice induced protective neutralizing antibody titers.

Notes: Mice immunized with the various formulations were challenged with a low pathogenic influenza virus 63 days after the initial vaccination. One week before the challenge, all vaccine formulations induced high neutralizing antibody titers consistent with previous work and were suggested to be protective. Error bars represent the standard error of the mean. Different letters indicate statistical significance among treatments. $P \leq 0.0001$.

Abbreviations: ID₅₀, sera dilution that inhibits 50% of pseudovirus infectivity; H5₃, H5 hemagglutinin trimer; sH5₃, soluble H5 hemagglutinin trimer.

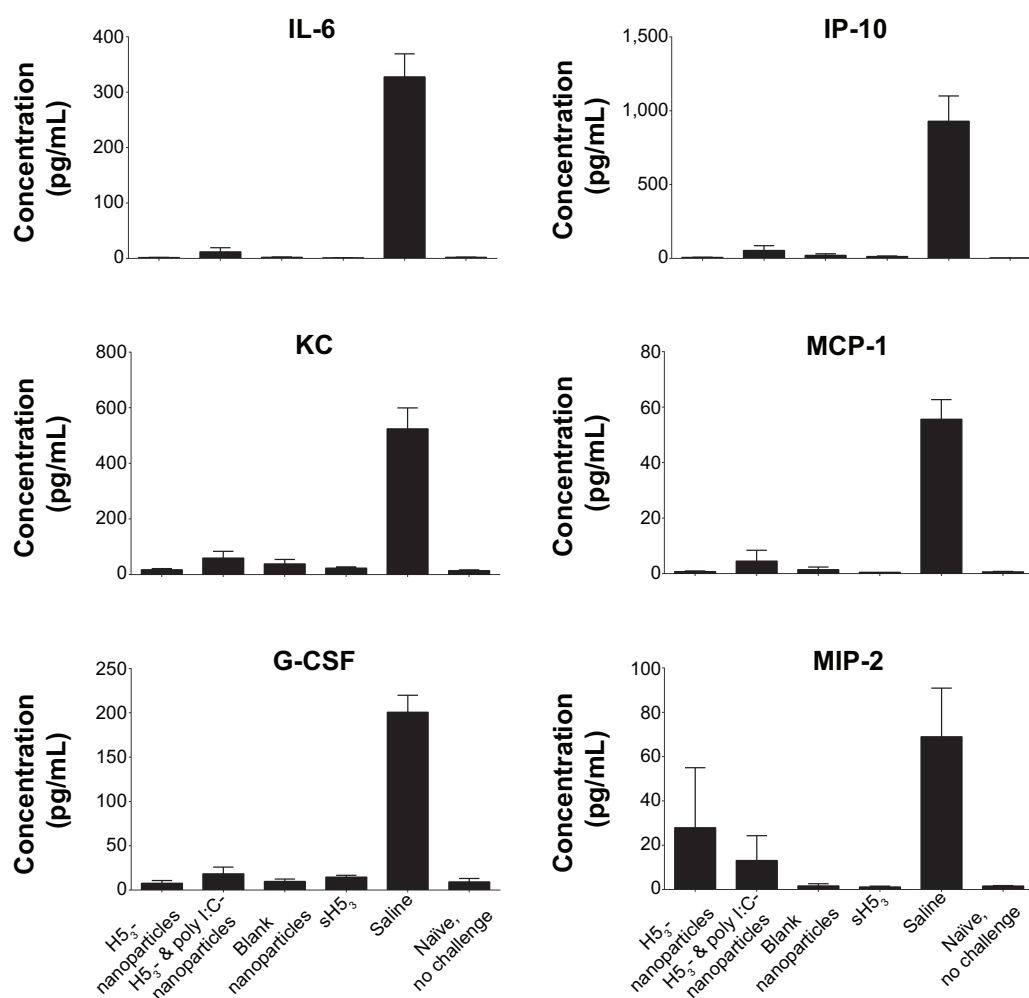


Figure S2 Low levels of inflammatory cytokines were present in immunized mice postchallenge.

Notes: Saline-administered mouse lungs had significant increases in inflammatory cytokines 3 days postchallenge. In contrast, the lungs of H5₃-immunized mice had low levels of inflammatory cytokines similar to healthy, noninfected mice.

Abbreviations: H5₃, H5 hemagglutinin trimer; sH5₃, soluble H5 hemagglutinin trimer.

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