

RESEARCH ARTICLE



Antibody responses against heterologous H5N1 strains for an MF59-adjuvanted cell culture–derived H5N1 (aH5N1c) influenza vaccine in adults and older adults

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ABSTRACT

MF59-adjuvanted H5N1, cell culture-derived inactivated influenza vaccine (aH5N1c, AUDENZ[®], Seqirus) is available for persons 6 months of age and older. During a pandemic, lack of preexisting immunity to novel influenza strains increases morbidity and mortality. This study examined the potential for an adjuvanted vaccine to provide cross-protection to novel viruses. Two similarly designed studies involving separate cohorts aged 18–64 and ≥65 y assessed immune responses to five heterologous H5N1 influenza strains elicited by two 7.5 µg doses of aH5N1c given 3 weeks apart. Geometric mean titers (GMT) on Days 1 and 43 and Day 43/Day 1 geometric mean ratios (GMRs) were determined with hemagglutination inhibition (HI) and microneutralization (MN). Rates of seroconversion (SC) and percentages of subjects with HI and MN ≥ 1:40 were determined. Significant increases in GMTs were observed on Day 43 after vaccination for all 5 heterologous strains in all ages tested. SC rates were 28–55% and 17–46% among those aged 18–64 and ≥65 y, respectively. MN ≥ 1:40 was observed in 38–100% of younger and 37–97% of older subjects, and HI ≥ 1:40 was achieved by 28–64% of subjects aged 18–64 y and by 17–57% of subjects aged ≥65 y. A SC rate ≥40% (97.5% CI) was met for two heterologous strains tested in adults aged 18–64 y. In adults aged 18–64 and ≥65 y, two 7.5 µg doses of aH5N1c demonstrated increased immunogenicity from baseline against five heterologous H5N1 strains, illustrating the potential for aH5N1c to provide cross-protection against other H5N1 strains.

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Introduction

The COVID-19 pandemic has highlighted the tremendous burden pandemics can place on individuals and society. Likewise, influenza A pandemics involving novel influenza A viruses can dramatically increase deaths and hospitalizations.^{1–3} The potential for morbidity and mortality is higher with pandemic than seasonal influenza because the population has little or no preexisting immunity. During a non-pandemic flu season, children and older adults (i.e., those >65 y of age) tend to be most vulnerable to complications associated with influenza. However, during an influenza pandemic, the risk of severe influenza-associated complications for healthy adults may be similar to that of young children and older adults.³ Vaccines, the main prophylactic measure against pandemic influenza, thus have an important role in pandemic preparedness plans worldwide.^{2–4} Pandemic influenza strains result from an antigenic shift and may include hemagglutinin (HA) subtypes of either avian origin such as H5, H7, and H9 or swine variants of HA subtypes H1, H2, and H3 that have further acquired adaptive mutations to become infectious from human to human.^{5,6} Among these subtypes, the highly pathogenic avian influenza (HPAI) H5N1 subtype and its genetic reassortants (including H5N2, H5N5, H5N6, and H5N8) are a global threat not only to poultry and wildlife but also to poultry workers and the general population because these viruses can cross the animal-human barrier and the human population lacks preexisting immunity to H5 viruses. Most human cases reported to date

have resulted from exposure to poultry or to environments contaminated by poultry-borne viruses.^{7–9} Based on the latest World Health Organization (WHO) report of human infection with avian influenza H5 virus, from January 2003 to 25 November 2022, there have been 868 cases of human infection with avian influenza A(H5N1) virus reported from 21 countries. Of these 868 cases, 457 were fatal (case fatality rate of 53%).¹⁰

According to the US Department of Human and Health Services (DHHS) and the Department of Homeland Security, a vaccine specific to an emerging pandemic influenza strain should be available to vaccinate the entire US population within 6 months of a declaration of a pandemic.^{2,4} However, time is a limiting factor in the process to develop, approve, and produce a pandemic-specific vaccine in large quantities, especially when using egg-based production methods, which require very large numbers of fertilized eggs. Hence, these US agencies prioritize maintenance of pandemic influenza vaccine stockpiles, which can be used to vaccinate front-line personnel, including healthcare workers, first responders, and the military.⁴ Given the unpredictable nature of pandemic viruses, a stockpiled vaccine that can provide broad heterologous coverage would represent a major asset to pandemic control efforts. Such a vaccine would offer enhanced protection until an antigenically matched vaccine could be manufactured and distributed.^{11,12}

The adjuvant MF59 (Seqirus Inc., Boston, MA, USA) is an oil-in-water emulsion stabilized by Tween 80 and Span 85 that

enhances differentiation of immune cells into antigen-presenting cells and also promotes macrophage recruitment, antigen uptake, lymph node migration, T-cell activation, and B-cell expansion.^{13–20} MF59 adjuvant is present in egg propagated trivalent and quadrivalent vaccines licensed for immunization against seasonal influenza in adults ≥ 65 y, a recently licensed cell culture-derived H5N1 vaccine, and two versions of egg propagated H1N1 vaccine licensed during the 2009 pandemic, all for persons 6 months of age or older. MF59 promotes production of cross-reactive antibodies against antigenically different influenza strains and has a well-established safety profile.^{21–25} In a recent pair of studies with identical designs, aH5N1c demonstrated a robust immune response to the homologous virus strain in separate adult populations aged 18–64 y and ≥ 65 y.²⁶ This report examines the immunogenicity of aH5N1c against heterologous strains of H5N1 in the same two study populations.

Materials and methods

Study design

Two phase 2, randomized, observer-blind, multicenter studies involving adults 18–64 y (NCT01776541) and older adults aged ≥ 65 y (NCT01766921) were conducted in Australia, New Zealand, the United States, and Thailand. The design, objectives, and endpoints were identical for both studies, as previously described.²⁶ The studies were conducted in compliance with Good Clinical Practices guidelines and the Declaration of Helsinki, and study protocols were approved by the Ethics Review Committees of the participating centers. Written informed consent was obtained from subjects before enrollment.

In a pre-defined exploratory analysis, hemagglutination inhibition (HI) and microneutralization (MN) antibody responses against five H5N1 heterologous influenza strains were evaluated in a randomly selected subset of subjects from the full-dose treatment arm from these two phase 2 trials.

Vaccine administration

The aH5N1c vaccine contained A/turkey/Turkey/1/05 (H5N1)-like strain (NIBRG-23) antigen (Seqirus Inc., NC, USA; f/k/a Novartis Influenza Vaccines GmbH, Marburg, Germany). Subjects were randomized at a 1:1 ratio to receive two vaccinations with either full-dose (7.5 μ g of hemagglutinin antigen per dose, 0.25 mL MF59) or a half-dose (3.75 μ g of hemagglutinin antigen per dose, 0.125 mL MF59). Vaccines were administered on Day 1 and Day 22 as single intramuscular injections in the non-dominant arm. Only subjects who received the 7.5 μ g antigen dose of aH5N1c were included in the heterologous analysis.

Participants

As described previously, adults aged 18–64 y and those aged ≥ 65 y were enrolled in separate studies. The main exclusion criteria were the presence of serious chronic or progressive disease; pregnancy or breastfeeding; prior receipt of any H5N1 vaccine; receipt of any other influenza vaccines within 60 d

prior to enrollment; a body mass index ≥ 35 kg/m², body temperature $\geq 38.0^\circ\text{C}$, and/or any acute illness within 3 d of receiving study vaccines. The present analysis of heterologous responses includes a prespecified subset of subjects receiving the full dose of study vaccine; heterologous responses to the half dose were not evaluated.

Endpoints

Antibody responses against the homologous H5N1 strain A/turkey/Turkey/1/2005 and the following heterologous strains were measured by HI and MN assays: A/Anhui/2005 CC Ab (Clade 2.3.4), A/Egypt/2010 CC Ab (Clade 2.2.1), A/Hubei/2010 CC Ab (Clade 2.3.2.1), A/Indonesia/2005 CC Ab (Clade 2.1.3), and A/Vietnam/1203/2004 CC Ab (Clade 1), selected to represent various genetic clades and subclades from WHO-identified H5N1 viruses of concern. Measures of immunogenicity included geometric mean titers (GMTs) on Days 1 and 43, Day 43/Day 1 geometric mean ratio (GMR) of HI and MN titers, the percentage of subjects with HI $\geq 1:40$ or MN $\geq 1:40$ on Days 1 and 43, and percentage of subjects achieving seroconversion (defined as HI $\geq 1:40$ for subjects negative at baseline [HI < 1:10] or a minimum 4-fold increase in HI titer for subjects positive at baseline [HI $\geq 1:10$] or a ≥ 4 -fold rise in MN titers on Day 43).

Statistical methods

Statistical methodology of the primary studies has been previously described.²⁶ The percentages of subjects with HI titers $\geq 1:40$ and achieving seroconversion were calculated along with the associated 97.5% Clopper-Pearson confidence intervals (CIs). Although the Center for Biologics Evaluation and Research (CBER) licensure criteria define lower limits of 95% CIs, 97.5% CIs were calculated because two vaccine formulations were assessed, and the 0.05 alpha was distributed across tests. GMTs, GMRs, and the associated 2-sided 95% CIs were calculated using analysis of covariance (ANCOVA) with factors for baseline titer and study center applied on log₁₀-transformed values. For the homologous strain, similar methods were used.

Results

Study population

Of subjects enrolled to receive the full dose aH5N1c in the two primary studies, 488 were 18–64 y of age (mean age 39 y) and 700 were ≥ 65 y (mean age 71 y).²⁶ Among both younger and older adults, the majority were white and female. More older than younger adults had an influenza vaccination within the prior 12 months, and baseline characteristics were similar between the age groups (Table 1). HI antibody responses, for subjects who had blood drawn/serum availability, to the homologous strain were tested in 478 adults aged 18–64 y and 693 adults aged ≥ 65 y in the primary studies,²⁶ and MN titers against the homologous strain and HI and MN titers against heterologous strains were measured in 69 and 35 younger and older adults, respectively, in this exploratory analysis. The sample size for the heterologous strain assays was based on historical precedents for influenza descriptive statistics.

Table 1. Study population demographics.

	Age 18–64 y (n = 488)	Age ≥65 y (n = 700)
Age, mean ± SD (y)	39.0 ± 13.7	71.2 ± 5.1
Male, n (%)	203 (42)	293 (42)
Weight, mean ± SD (kg)	74.1 ± 15.1	71.0 ± 15.9
Height, mean ± SD (cm)	167.8 ± 10.5	164.1 ± 10.7
BMI, mean ± SD (kg/m ²)	26.2 ± 4.3	26.1 ± 4.0
Previous influenza vaccination, n (%)	118 (24)	429 (61)
Influenza vaccination within 12 months prior to study, n (%)	98 (20)	388 (55)
Race, n (%)		
White	291 (60)	445 (64)
Black	97 (20)	10 (1)
Asian	93 (19)	240 (34)
Native American	2 (<1)	2 (<1)
Other	5 (1)	3 (<1)
Hispanic ethnicity, n (%)	107 (22)	17 (2)

Immunogenicity

As shown in Table 2, on Day 43 (3 weeks after the second vaccination), HI GMT increased 41-fold vs Day 1 for the homologous strain and by 2- to 12-fold against all five heterologous strains in subjects aged 18–64 y. As measured with MN in the same population, homologous GMT was 61 times higher and heterologous GMTs were 5–34 times higher on Day 43 vs Day 1. In adults aged ≥65 y, homologous GMTs increased 16- and 23-fold as measured with HI and MN assays, respectively. Heterologous HI GMTs increased by 2- to 5-fold and MN GMTs by 4- to 12-fold.

HI ≥ 1:40 was achieved by 28% to 64% of subjects aged 18–64 y and by 17% to 57% of subjects aged ≥65 y (Figure 1). MN ≥ 1:40 was observed in 38% to 100% of younger and 37% to 97% of older subjects (Figure 2).

HI seroconversion rates against heterologous strains ranged from 28% to 55% among adults 18–64 y and 17% to 46% among those ≥65 y of age. The lower bound of the 97.5% CI of the seroconversion rate in adults aged 18–64 y was ≥40% for the Egypt and Hubei heterologous strains (Figure 3(a,b)). The percentage of subjects with a ≥ 4-fold increase in MN titers against heterologous strains ranged from 32% to 88% in

subjects aged 18–64 y and 26% to 74% for older adults. Most younger adults achieved a ≥ 4-fold increase in MN against all but the Vietnam strain, whereas among older adults, the majority achieved this threshold only against the Egypt strain (Figure 3(c,d)).

Discussion

This exploratory analysis confirmed that aH5N1c not only elicits a robust immune response against the homologous strain but also against heterologous strains of the H5N1 pandemic influenza virus in adults younger and older than 65 y. CBER criteria (97.5% CI of the seroconversion rate ≥40%) were met for the Egypt/2010 and Hubei/2010 strains in adults aged 18–64 y. Thus, this vaccine could provide essential interim coverage to a population – particularly those 10–40 y of age where H5N1 morbidity and mortality is highest – threatened by an H5N1 pandemic while strain-specific vaccine was being developed and manufactured.

Influenza pandemics have historically been characterized by shifts in the virus subtype, successive pandemic waves, higher transmissibility compared with seasonal influenza,

Table 2. HI and MN antibody responses (GMT and GMR) against H5N1 at Day 1 and Day 43 by age group.

	HI			MN		
	GMT Day 1 (95% CI)	GMT Day 43 (95% CI)	GMR Day 43/Day 1 (95% CI)	GMT Day 1 (95% CI)	GMT Day 43 (95% CI)	GMR Day 43/Day 1 (95% CI)
Age 18–64 y						
Homologous	n = 478	n = 451	n = 451	n = 69		
Turkey/2005	6.11 (5.78–6.46) ^a	250 (208–302) ^a	41 (34–49) ^a	6.7 (5.62–7.98)	410 (313–537)	61 (44–86)
Heterologous		n = 69		n = 69		
Anhui/2005	5.02 (4.87–5.18)	11 (6.64–17)	2.09 (1.29–3.37)	16 (12–21)	156 (131–186)	9.93 (7.47–13)
Egypt/2010	5.87 (4.51–7.63)	39 (22–71)	6.52 (3.62–12)	15 (11–22)	522 (398–685)	34 (23–49)
Hubei/2010	7.12 (5.01–10)	56 (31–102)	7.3 (4.01–13)	7.2 (6–8.63)	51 (41–63)	7.07 (5.56–8.98)
Indonesia/2005	5.02 (4.87–5.18)	16 (9.17–28)	3.12 (1.81–5.39)	8.79 (7–11)	114 (89–146)	13 (9.42–18)
Vietnam/1203/2004	5.97 (4.41–8.06)	44 (24–81)	6.99 (3.84–13)	5.59 (4.98–6.27)	27 (21–34)	4.76 (3.74–6.06)
Age ≥65 y						
Homologous	n = 693	n = 673	n = 673	n = 35		
Turkey/2005	8.29 (7.57–9.08) ^a	129 (110–152) ^a	16 (13–19) ^a	7.49 (5.87–9.56)	169 (107–268)	23 (13–39)
Heterologous		n = 35		n = 35		
Anhui/2005	5 (5–5)	8.53 (5.92–12)	1.71 (1.18–2.46)	28 (20–41)	128 (99–165)	4.52 (2.93–6.97)
Egypt/2010	6.86 (4.94–9.53)	27 (15–47)	3.88 (2.27–6.64)	25 (15–41)	303 (206–447)	12 (6.69–22)
Hubei/2010	7.43 (5.28–10)	36 (20–64)	4.83 (2.8–8.32)	11 (7.98–16)	42 (31–58)	3.75 (2.6–5.4)
Indonesia/2005	5 (5–5)	12 (8.02–18)	2.39 (1.6–3.56)	18 (13–27)	87 (63–120)	4.77 (3.1–7.34)
Vietnam/1203/2004	7.81 (5.24–12)	34 (19–63)	4.41 (2.57–7.57)	7.32 (5.39–9.95)	27 (19–39)	3.68 (2.62–5.17)

Abbreviations: CI, confidence interval; GMR, geometric mean ratio; GMT, geometric mean titer; HI, hemagglutination inhibition; MN, microneutralization.

^a97.5% CI.

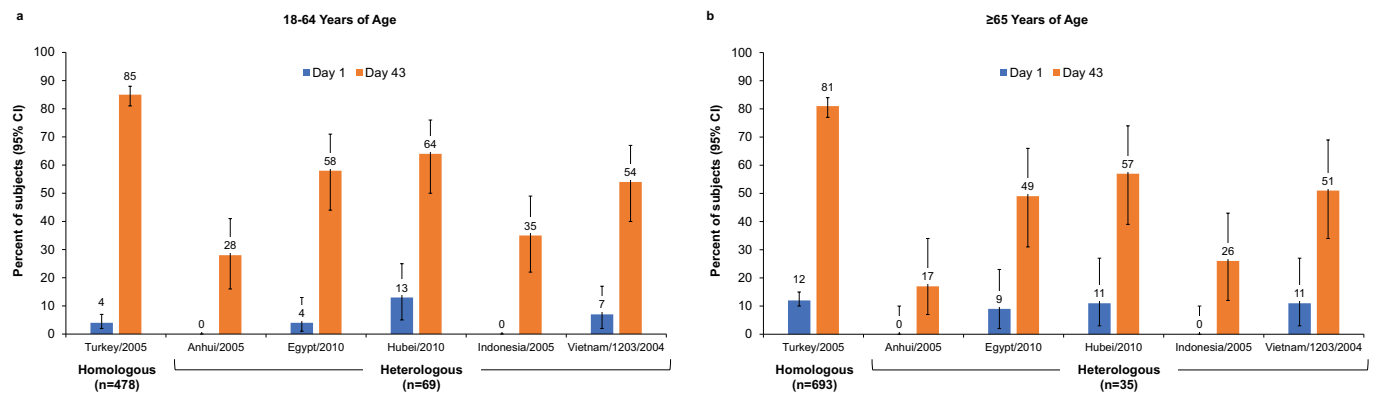


Figure 1. Percentage of subjects with hemagglutination inhibition (HI) titers $\geq 1:40$ in adults aged 18–64 y (a) and aged ≥ 65 y (b). Error bars represent the 95% confidence interval (CI) for heterologous strains and the 97.5% CI for the homologous strain.

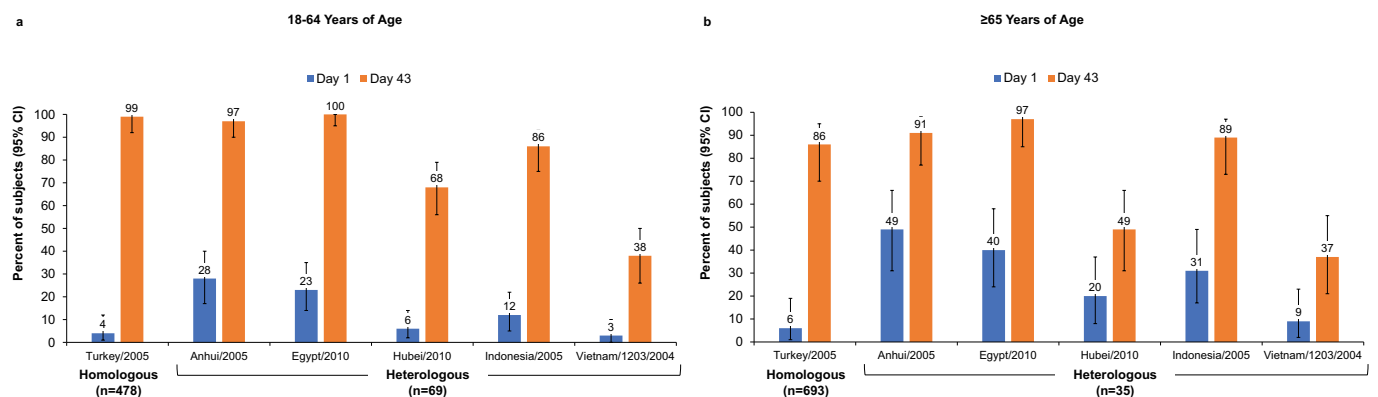


Figure 2. Percentage of subjects with microneutralization (MN) titers $\geq 1:40$ in adults aged 18–64 y (a) and aged ≥ 65 y (b). Error bars represent the 95% confidence interval (CI) for heterologous strains and the 97.5% CI for the homologous strain.

and variable impact in different geographic regions, as well as increased morbidity and mortality in older children and adults younger than 65 y – the population typically most resistant to seasonal influenza complications.³ The vulnerability of younger adults to pandemic influenza strains is of special concern, as this age group is most likely to be on the front lines of the pandemic response, providing health care and other essential services.⁴ The cross-reactivity against five antigenic variants shown by aH5N1c in this analysis suggests this vaccine could provide these essential workers with cross-protection during the period between the start of a pandemic and distribution of an antigenically matched vaccine.

Millions of fertilized eggs are needed to produce a vaccine, and efficiency of this method is often low, requiring one or more eggs for each dose of the vaccine produced. These constraints could pose challenges during a highly pathogenic avian influenza outbreak or pandemic if either egg quantity or quality become compromised. Cell culture manufacturing methods offer efficient vaccine production that minimizes dependency on the egg supply and may provide increased efficacy relative to egg-derived vaccines.^{27–32} In addition, enhanced immunogenicity with MF59 adjuvant against heterologous strains, when added to both seasonal and pandemic influenza vaccines, is well documented in children and adults.^{33–37}

Observed differences between MN and HI titers in this study may be due to amino acid substitutions found in the

receptor-binding site of the HA molecule of influenza A strains, which can alter the results of HI assays when cell culture-derived vaccines are evaluated.^{38–40} The MN assay is not affected by HI mutations because MN measures the concentration of antibodies needed to prevent infection of a eukaryotic cell.^{41,42} It is possible that the MN assay may represent a more mechanistically relevant estimation of antibody-mediated protection than HI.^{43,44} The choice of erythrocyte species can influence the HI titer, which may also explain the lack of correlation between HI and MN for some results in this study.⁴⁵ The discrepancy in sample size between the HI and MN analysis groups could also have contributed to differences in titers. However, overall, the trends were consistent between the two assays (i.e., when HI increased, so did MN).

This study was limited by the small number of sera samples tested from the subgroup in which heterologous responses were assessed. Nevertheless, responses were robust, with CBER criteria being met for two of five strains despite the small sample size. The five heterologous strains tested represented a broad spectrum of possible H5N1 variants, although these were previously identified strains, and it cannot be predicted whether the responses to new strains would be similar. A limitation of the studies included in this analysis is that antibodies with H5-specific Fc-mediated effector functions were not determined. Vaccination with adjuvanted H7N9 and H5N1 vaccines

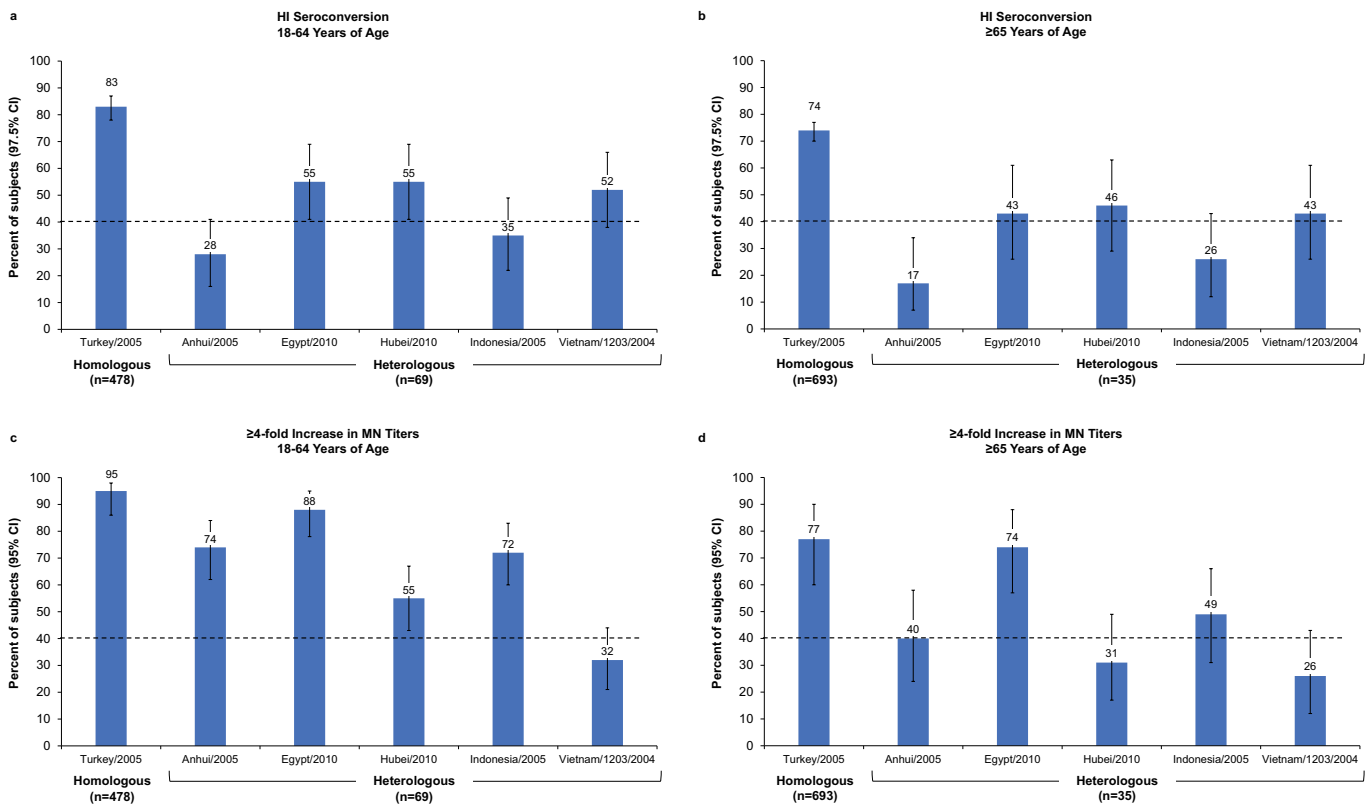


Figure 3. Percentage of subjects with seroconversion (defined as hemagglutination inhibition [HI] $\geq 1:40$ for subjects negative at baseline [HI $< 1:10$] or a minimum 4-fold increase in HI titer for subjects positive at baseline [HI $\geq 1:10$]) on Day 43) or a ≥ 4 -fold increase in microneutralization (MN) titers. (a, b) Seroconversion in adults aged 18–64 y (a) and ≥ 65 y (b). Error bars represent the 97.5% confidence interval (CI). (c, d) at least 4-fold increase in MN titers in adults aged 18–64 y (c) and ≥ 65 y (d). Error bars represent the 95% CI.

has been shown to induce antibody-dependent cellular cytotoxicity (ADCC)-mediating antibodies. Protection from influenza may require Fc-mediated antibody functions in addition to neutralizing antibodies. Additional studies are necessary to understand how and to what extent ADCC- and antibody-dependent phagocytosis (ADP)-mediating antibodies contribute to protection from influenza, as these antibodies may also have an immunopathological role.

In conclusion, in adults younger than 65 y and those 65 y and older, two 7.5 μg doses of aH5N1c, administered three weeks apart, demonstrated increased immunogenicity from baseline against multiple heterologous H5N1 strains, of five separate genetic clades or subclades. These findings illustrate the potential for the aH5N1c vaccine to provide cross-protection against other H5N1 strains during a pandemic and provide support for future research in this area.

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Author contributions

All authors participated in either the conception and design of the study, acquisition of data and/or analysis and interpretation of data. In addition, all authors participated in drafting and critically revising the article for important intellectual content, and all authors provided final approval of the submitted version.

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