Inactivated and live-attenuated seasonal influenza vaccines boost broadly neutralizing antibodies in children

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In brief
Yegorov et al. evaluate vaccine-elicited broadly neutralizing antibody (bNAb) responses against group 1 influenza A viruses in children. Repeated seasonal immunization of children against influenza boosts bNAb titers. Inactivated and live attenuated vaccine formulations are similarly effective at boosting bNAbs, with the magnitude of boosting inversely correlated with age.

Highlights
- Repeated inactivated influenza vaccination boosts bNAbs
- Inactivated and live attenuated vaccines are similarly efficient at boosting bNAbs
- The magnitude of IIV and LAIV vaccine-elicited bNAb boosting declines with age
Inactivated and live-attenuated seasonal influenza vaccines boost broadly neutralizing antibodies in children

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https://doi.org/10.1016/j.xcrm.2022.100509

SUMMARY

The induction of broadly neutralizing antibodies (bNAbs) that target the hemagglutinin stalk domain is a promising strategy for the development of “universal” influenza virus vaccines. bNAbs can be boosted in adults by sequential exposure to heterosubtypic viruses through natural infection or vaccination. However, little is known about if or how bNAbs are induced by vaccination in more immunologically naive children. Here, we describe the impact of repeated seasonal influenza vaccination and vaccine type on induction of bNAbs against group 1 influenza viruses in a pediatric cohort enrolled in randomized controlled trials of seasonal influenza vaccination. Repeated seasonal vaccination results in significant boosting of a durable bNAb response. Boosting of serological bNAb titers is comparable within inactivated and live attenuated (LAIV) vaccinees and declines with age. These data provide insights into vaccine-elicited bNAb induction in children, which have important implications for the design of universal influenza vaccine modalities in this critical population.

INTRODUCTION

Despite decades of study and the accessibility of seasonal vaccination, influenza remains an important public health concern, largely owing to its ability to escape immunity and cause seasonal epidemics and pandemics.1–3 Seasonal vaccines, the mainstay of influenza prevention, have often shown modest real-world effectiveness due to suboptimal immunogenicity, alongside rapid and unpredictable changes in circulating virus strains.1–3 Globally, several different seasonal influenza vaccine formulations are approved, including intramuscular injection of inactivated virus (IIV) or mucosal administration of replication-competent live attenuated influenza vaccine (LAIV).1 Although influenza vaccines are often considered equally efficacious, there are insufficient data on the impact of repeated vaccination and vaccine type on the breadth of influenza-virus-directed immunity in specific demographic groups, especially children, who are a major source of influenza transmission and, alongside the elderly, are at heightened risk for influenza-related hospitalizations and death.4,5

The bulk of influenza-vaccine-elicited antibodies (Abs) target the viral membrane surface proteins hemagglutinin (HA) and, to a lesser extent, neuraminidase (NA). Of the HA-reactive Abs, most bind the “immunodominant” globular HA head domain in a strain-specific manner, blocking the interaction between the virus and cell surface sialic acid, while a minority of HA-reactive Abs bind the sub-dominant, but highly conserved HA stalk domain; a subset of these Abs is considered broadly neutralizing (bNAbs). HA stalk-reactive bNAbs have emerged as a promising strategy for the development of a “universal” influenza vaccine.6,7 In adults, we and others have shown that bNAbs are selectively boosted by exposure to pandemic/heterosubtypic HA subtypes (via either infection or vaccination).8–10 bNAb titers also increase over time owing to an accumulation of lifetime exposures to influenza virus selecting for cross-reactive B cell clone maintenance and expansion.11,12 A recent clinical trial demonstrated that chimeric hemagglutinin vaccines delivered in live attenuated and inactivated formulations could successfully boost bNAb titers in adults.6,7 However, little is known about how bNAbs develop in children and how vaccination impacts bNAb development—knowledge that will have important implications for understanding the nature of vaccine-mediated immunity against influenza, especially in the context of exposure to novel strains.1,13,14
Therefore, we explored the effects of seasonal influenza vaccination and vaccine formulation on bNAb induction in participants of a cluster randomized controlled trial (cRCT) designed to compare the community-level protection mediated by IIV and LAIV vaccination of children. We found that titers of neutralizing bNAbs were enhanced by repeated seasonal vaccination and that bNAb levels were boosted similarly by LAIV and IIV. As expected, LAIV and IIV effects on bNAb titers were more pronounced in the mucosa and blood, respectively, and the magnitude of vaccine-induced bNAb responses in the serum declined with age.

**RESULTS**

**Study participants**
The effects of repeated seasonal IIV vaccination on the induction of bNAbs were studied using serum samples from 68 RCT participants (median age 9.0 years) (Table S1), who either received IIV (37 vaccinees) or served as controls (31 controls) for each of the three seasons (2008/09 to 2010/11) captured by the cRCT (Figure 1A and Table 1) and for whom paired samples were available. The effects of vaccine type on bNAb elicitation were subsequently studied using serum and mucosal samples from 72 RCT participants (median age 11.0 years) (Table S1), who received either the inactivated vaccine (IIV, n = 35) or the live attenuated influenza vaccine (LAIV, n = 37) (Figure 1B and Table 1) and for whom paired samples were available.

**Effects of repeated seasonal vaccination on serological bNAb responses**
To determine the effect of repeated seasonal IIV vaccination of children on the induction of bNAbs, microneutralization (MNT) assays were performed on the 2008/09 and 2012/13
globulin G (IgG) and IgA and bNAb (anti-cH5/1 N3) MNT50 titers formulation (IIV versus LAIV) influences the magnitude of serolog-

tion mediated by group 1 HA stalk-binding bNAbs.9,10,13,14 After 

N3, a chimeric virus previously validated to quantify neutraliza-

pre-vaccination sera (corresponding to the beginning of the 

cRCT and 1 year after its conclusion, respectively) using ch5/1 

N3, a chimeric virus previously validated to quantify neutralization 

mediated by group 1 HA stalk-binding bNAbs.5,10,13,14 After 

three vaccination seasons, higher MNT50 titers were evident in 

vaccinates compared with controls (geometric mean fold change 

(GMFC) = 2.65 versus 1.43, p < 0.001) (Figures 2A, 2C,S1, and 

S3). The boost in anti-Cal09 H1N1 IgG and HAI was significantly 

higher in the IIV vaccinees compared with the LAIV vaccinees 

(S2; Table S2); 43% (16/37) of vaccinees had at least a 4-fold in-

crease of MNT50 compared with the 13% (4/31) seroconversion 

among controls. bNAbs that bind to the receptor-binding site 

(RBS) of HA have also been reported.15 To determine whether 

repeated vaccination boosted titers of this antibody class, we 

did hemagglutination inhibition (HAI) assays using the cH5/1 

N3 virus employed in the MNT assay. No increase in HAI-positive 

vaccinees compared with controls (geometric mean fold change 

(GMFC = 4.59 versus 2.0, p = 0.017 for IgG, and GMFC = 3.03 

versus 1.62, p = 0.007 for HAI) (Figures 4D and 4F). There was 

no difference in boosting magnitude between the two groups 

for Cal-09 H1N1 IgA responses, though the absolute post-boost 

HAI titers were higher for IIV vaccinees (Figures 4E and 4F). The 

magnitude of vaccine-elicited Cal/09 H1N1 serum IgA increase 

was inversely correlated with age in all vaccinees (r = −0.379, 

p = 0.001) (Figure 4H). Boosting of Cal/09 H1N1 HAI was 

also inversely correlated with age for IV vaccinees (r = −0.359, 

p = 0.034) (Figure 4I). Thus, the boost in functional antibody re-

sponses against vaccine antigen appears more pronounced in 

IVV and exhibits some age dependence.

Effects of vaccine type on mucosal IgG and IgA responses
Vaccine-elicited mucosal immune responses are critical for pro-
tection against influenza. Therefore, we next assessed the anti-
body profiles in nasal swabs to determine the impact of vaccine 
type on mucosal immunity. Overall, vaccination tended to in-
duce mucosal bNAb titers, without significant differences be-
tween vaccine types (Figures 5A, 5B, 5E, and 5F). Anti-stalk 
IgG was significantly elevated, with the largest effect size in 
LAIV vaccinees (GMFC = 3.17, p < 0.001) (Figures 5A and 5E). 
Notably, alongside an apparent increase in IgG, there was a 
decrease in IgA titers against the Cal/09 H1N1 vaccine antigen 
in LAIV vaccinees, as reported previously.15 No significant corre-
lations were observed between vaccine-mediated boosting of 
mucosal Abs and age (Figures 5I–5L). Altogether, LAIV and IVV 
boosted mucosal IgG and IgA bNAbs to a similar extent.

Relationship between vaccine-elicited serological and 
mucosal antibody response
Whether and how vaccine-elicited immunity correlates 
between the systemic and mucosal compartments is a topic of 
debate.20,21 Therefore, we next explored the relationship

Table 1. Description of influenza strains used in the standard-of-care trivalent vaccines administered over the study period

| Season     | Strain type | Sub-type | Strain name             |
|------------|-------------|----------|-------------------------|
| Analysis I seasons |
| 2008/09    | A           | H1N1     | Brisbane/59/2007        |
| 2008/09    | A           | H3N2     | Brisbane/10/2007        |
| 2008/09    | B           | Yamagata | Florida/4/2006          |
| 2009/10    | A           | H1N1     | Brisbane/59/2007        |
| 2009/10    | A           | H3N2     | Brisbane/10/2007        |
| 2009/10    | B           | Victoria | Brisbane/60/2008        |
| 2010/11    | A           | H1N1     | California/7/2009       |
| 2010/11    | A           | H3N2     | Perth/16/2009           |
| 2010/11    | B           | Victoria | Brisbane/60/2008        |
| Analysis II seasons |
| 2014/15    | A           | H1N1 (pdm09) | California/7/2009   |
| 2014/15    | A           | H3N2     | Texas/50/2012           |
| 2014/15    | B           | Yamagata | Massachusetts/2/2012    |

Effects of vaccine type on serological bNAb responses
IIV and LAIV vaccines are known to elicit distinct immune responses.17–19 Thus, we set out to determine whether vaccine formulation (IIV versus LAIV) influences the magnitude of serological vaccine-induced bNAb responses in children. There was a significant elevation of HA stalk-binding (anti-ch6/1) immunoglobulin G (IgG) and IgA and bNAb (anti-ch5/1 N3) MNT50 titers at 1 month after vaccination (GMFC = 2.71, p < 0.001; 1.51, p = 0.004; 1.48, p = 0.002, respectively) (Figures 3A–3C, Table S3); this elevation was more pronounced in IVV vaccinees than in LAIV vaccinees, although the response magnitude was not significantly different between the two groups (Figures 3D and 3F). The magnitude of vaccine-elicited boosting of both anti-HA stalk IgG and MNT50 (r = −0.238, p = 0.045 and r = −0.333, p = 0.004, respectively), but not anti-HA stalk IgA titers, declined with age (Figures 3G–3I). There were no significant differences in vaccination history between the IVV and LAIV groups (Table S4). Altogether, these data show that in young children, both IVV and LAIV boost serum bNAbs in an age-dependent manner.
between the magnitude of vaccine-induced bNAb boost in the serum versus the nasal mucosa. In unstratified analysis, no correlation was observed between the serological and mucosal antibody responses (Figures 6A–6D, left panels). Upon stratification by vaccine type, there was no correlation between serum and mucosal responses for bNAbs (Figures 6A and 6B, right panels). However, the magnitude of change in vaccine Cal/09 H1N1-reactive IgG, but not Cal/09 H1N1-reactive IgA, positively correlated between serum and mucosa in IIV vaccinees only ($r = 0.442$, $p = 0.011$) (Figures 6C and 6D, right panels). Thus, bNAb responses appear to be compartment specific.

**DISCUSSION**

Notwithstanding the recent global decline in seasonal influenza incidence as a consequence of public health measures, 

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**Figure 2. Effects of seasonal vaccination of bNAb titers in children**

(A) Anti-HA stalk MNT$_{50}$ titers at baseline (year 2008) and after three cRCT vaccination seasons (year 2012).
(B) Serum HAI activity against the Cal/09 virus assessed at baseline (year 2009) and after three cRCT vaccination seasons (year 2012).
(C) Log$_2$-transformed ratios of MNT$_{50}$ titers after repeated vaccination (year 2012) versus at baseline (year 2008).
(D) Log$_2$-transformed ratios of HAI titers after repeated vaccination (year 2012) versus at baseline (year 2009).
(E and F) Correlation plots of log$_2$-transformed MNT$_{50}$ (E) and HAI (F) titer ratios and participant ages at enrollment.

In (A and B), dots and brackets represent geometric means and 95% CIs, respectively; $p$ values indicate the statistical significance of the intra-individual difference between pre- and post-vaccination titers assessed by paired $t$ test. In (C and D), dots and brackets represent means and 95% CIs, respectively; $p$ values indicate the statistical significance of the difference between vaccinees and controls assessed by unpaired $t$ test. In (E and F), the Pearson coefficients ($r$) and their statistical significance are shown along with lines of best fit for vaccinees.
influenza remains an important public health concern owing to its inevitable seasonal reemergence as social distancing eases and the ever-present risk of future pandemics. Young children are a major driver of virus transmission, accounting for a significant proportion of influenza-related morbidity and mortality. Therefore, understanding how seasonal vaccination and varied vaccine formulations shape childhood immunity to influenza A viruses is of critical importance for effective influenza prevention.

Vaccine-elicited immune responses are tightly linked to history of exposure to virus, infection, and/or vaccination, and >50% of children have encountered influenza virus by 6 years of age. Previous studies have shown that vaccination of children with a pandemic H5 vaccine was capable of eliciting bNAbs—similar to what has been reported in adults. A major strength of our study is that it describes the impact of seasonal vaccination on bNAb induction in children whose exposure histories were carefully monitored within a cRCT framework, excluding the confounding effects of symptomatic infection while focusing specifically on the effects of immunization.

After three vaccine seasons, bNAb-mediated virus neutralization in IIV vaccinees increased by >2.5-fold, supporting the notion that seasonal vaccination facilitates sustained expansion of cross-reactive Ab repertoires. In contrast, although vaccine-elicited strain-specific HAI activity also rose over three seasons, it was not significantly different between the vaccinated and unvaccinated children at the conclusion of the study.

The magnitude of bNAb expansion induced by repeated IIV vaccination was inversely correlated with age, consistent with preferential boosting of Abs against the HA head domain that is known to occur in adults after seasonal influenza vaccination. Notably, a very modest bNAb elevation was observed in control subjects between 2008 and 2012, which is consistent
with the observation that HA stalk-reactive bNAbs tend to be polyreactive and that polyreactive Abs tend to increase with age.\textsuperscript{27} The rise in bNAbs in the control group also corresponded to the wave of 2009 H1N1 swine flu pandemic. Circulation of this virus in Canada began in the spring of 2009, outside of the usual influenza season and at a time when active surveillance was not being conducted in the context of this study. Exposure to this virus is well known to boost bNAbs, and we speculate that unrecognized infections may also partially explain this increase.\textsuperscript{28} In contrast, HAI titer increases observed in vaccinees and controls during the study period were very similar, and overall, these titers remained very low—below the typical seroprotection threshold of 40. Together, these data suggest that seasonally repeated vaccination boosted bNAbs in young children.

Importantly, IIV and LAIV were similarly effective at boosting bNAbs, in line with the equivalent efficacy offered by these two vaccine types in the cRCT.\textsuperscript{29} Furthermore, vaccine-elicited bNAb boosting was more pronounced in the blood of IIV vaccinees and in the mucosa of LAIV vaccinees, consistent with the respective vaccination routes and the results of previous studies comparing immune responses elicited by IIV and LAIV.\textsuperscript{30–32}

The role of mucosal immunity in protecting against respiratory virus infections has been somewhat enigmatic. On one hand, people with selective IgA deficiency have relatively modest increases in susceptibility to infections.\textsuperscript{33} Still, serum IgA has been reported to correlate with protection against influenza, and nasal IgA has been reported to contribute to protection, especially in individuals with low virus-specific HAI titers.\textsuperscript{34,35} The upper respiratory tract antibody repertoire is dominated by secretory IgA, while in the lower respiratory tract IgG is more abundant.\textsuperscript{7} We have previously shown that broadly neutralizing IgA is more potent than IgG.\textsuperscript{13} This is, at least in part, due to...
the glycosylation of IgA, whereby sialic acids present on the antibody mediate a second mode of binding that inhibits viral entry. We have also recently shown that the influenza virus-IgA immune complexes signal through the FcγRI receptor on neutrophils to stimulate NETosis and that neutrophil extracellular traps are capable of inactivating influenza virus. Therefore, robust mucosal immune response stimulation should be a priority for the development of more broad and efficacious influenza vaccines.

We did not observe a correlation between bNAb responses in the blood and nasal mucosa, suggesting an important contribution from in situ antibody generation at the mucosa. Interestingly, there was an inverse correlation between Cal/09 IgA titers and age at enrollment as well as anti-Cal/09 HAI titers in the IIV group. IIV recipients tended to have higher HAI titers against Cal/09 than LAIV recipients. Preexisting Abs can inhibit vaccine responses. Thus, the higher antibody titers in the IIV group might have inhibited IgG+ memory B cell stimulation and reentry into the germinal center and subsequent class switching to IgA. Furthermore, while the serum bNAb and Cal/09-specific IgA titers were similar, the titers of Cal/09-reactive IgG were approximately 2-fold higher compared with HA stalk-reactive IgG, consistent with the immunodominant nature of the HA head domain. Notably, mucosal titers of bNAbs and strain-specific Abs were similar, suggesting that mucosal responses may be less focused on immunodominant epitopes.
Conclusions

Our data provide insights into the effects of repeated seasonal influenza vaccination and vaccine type on bNAb induction in children. Seasonal influenza vaccines in children can induce bNAbs against influenza A both in blood and in respiratory mucosa, which has important ramifications for the selection of universal vaccine platforms that could be effectively deployed in this population. Specifically, our data suggest that the threshold for inducing bNAbs in children may be lower than in adults, for whom seasonal vaccines do a poor job at boosting bNAbs owing to immunodominance of the HA head domain.9,39 In addition, our data suggest that IIV and LAIV platforms might be equally suitable for delivery of universal influenza vaccines to pediatric cohorts. Future studies are needed to explore factors responsible

Figure 6. Relationship between mucosal and serological antibody titers
(A–D) Correlation plots of anti-HA stalk IgG (A) and IgA (B) and anti-Cal09 IgG (C) and IgA (D) log2-transformed post-/pre-vaccination titer ratios. The Pearson coefficients (r) and their statistical significance are shown along with lines of best fit for all participants (black) and IIV (red) versus LAIV (blue) vaccinees.
Limitations of the study
Our analyses were based on a relatively small selection of samples derived from the original cRCTs owing to sample availability. As the objective of the study was to specifically interrogate the impact of vaccination of bNAbs in children, those who were infected during the trial were excluded. Though the attack rates of influenza in these trials were very low (5%–10%), it is possible that the antibody responses of infected individuals might be different from those who were not infected. For feasibility reasons, we assessed only bNAbs against group 1 HAs. In the future, it would be important to examine group 2 bNAbs as well. Our mucosal analyses were performed using nasal swabs, which precluded conventional antibody titrations owing to the low antibody levels typical for these samples. However, we have shown previously that we are able to reliably measure differences in mucosal IgA using the methods employed herein. Finally, the protectiveness of bNAbs has been difficult to directly establish in vivo. Strain-specific Abs are much more potent than bNAbs and likely contribute most directly to protection in many instances. The protective effects of bNAbs are likely to be more apparent in seasons wherein there is a substantial mismatch between vaccine strains and circulating strains or in the context of pandemics. Indeed, bNAbs have been reported to correlate with protection against pandemic H1N1 independently of HA-positive Abs, and the elicitation of bNAbs during pandemics has been postulated as a mechanism for the extinction of seasonal pre-pandemic strains. Rigorous prospective cohort-based studies are needed to shed additional light on bNAb-mediated protection in humans.

STAR METHODS
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SUPPLEMENTAL INFORMATION
Supplemental content can be found online at https://doi.org/10.1016/j.xcrm.2022.100509.

ACKNOWLEDGMENTS
The authors would like to thank the cRCT participants and study teams. This work was supported, in part, by a CIHR Project Grant (M.S.M.), M.S.M. was supported, in part, by a CIHR New Investigator Award, an Ontario Early Researcher Award, and a Canada Research Chair in Viral Pandemics. H.D.S. was supported, in part, by an Ontario Graduate Scholarship. S.Y. was supported, in part, by an M.G. DeGroote Postdoctoral Fellowship.

AUTHOR CONTRIBUTIONS
Conceptualization, M.S.M and M.L.; investigation and formal analysis, M.S.M., S.Y., D.B.C., K.B.G., J.C.A., C.V., J.W., K.R., and H.D.S.; writing – original draft, M.S.M., S.Y., D.B.C., and K.B.G.; writing – review and editing, M.S.M., M.L., S.Y., D.B.C., K.B.G., J.C.A., C.V., J.W., K.R., and V.T.; funding acquisition, M.S.M. and M.L.

DECLARATION OF INTERESTS
The authors declare no competing interests.

Published: February 3, 2022
Revised: November 16, 2021
Accepted: January 7, 2022
Published: February 3, 2022

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# STAR★METHODS

## KEY RESOURCES TABLE

| REAGENT or RESOURCE | SOURCE | IDENTIFIER |
|---------------------|--------|------------|
| **Antibodies**      |        |            |
| Anti-Influenza A Antibody, nucleoprotein, clone A1, biotin-conjugated | EMB Millipore | Cat#MAB8257B |
| Goat Anti-Human IgA-HRP | SouthernBiotech | Cat#2050-05, RRID#AB_2687526 |
| Goat anti-Human IgG (H+L) Secondary Antibody, HRP | Invitrogen/Thermo Fisher | Cat#31410, RRID#AB_228269 |
| **Bacterial and virus strains** |        |            |
| ch5/1 N3 virus | Engineered using plasmids encoding 6 wild-type A/Puerto Rico/8/34 (PR8) strain segments and ch5/1 HA and N3 NA from A/Swine/Missouri/4296424/06 (Miss/06) virus<sup>9</sup> | N/A |
| A/California/04/2009 (Cal/09) H1N1 | Gift from Dr. Peter Palese, Icahn School of Medicine at Mount Sinai, NY | NCBI taxon ID#641501 |
| MAX Efficiency™ DH10Bac Competent Cells | Gibco/Thermo Fisher | Cat#10361012 |
| **Biological samples** |        |            |
| Participant sera | This study | N/A |
| Participant nasal swabs | This study | N/A |
| Chicken erythrocytes in Alsever’s solution | Canadian Food Inspection Agency (CFIA) | N/A |
| **Chemicals, peptides, and recombinant proteins** |        |            |
| ch6/1 hemagglutinin (HA) construct | Engineered using plasmids encoding A/Mallard/Sweden/81/02 H6 head domain and a A/Puerto Rico/8/1934 (PR8) group 1 stalk<sup>9</sup> | N/A |
| Dulbecco Modified Eagle Medium (DMEM) | Gibco/Thermo Fisher | Cat# 11965092 |
| Fetal bovine serum | Gibco/Thermo Fisher | Cat#12484028 |
| Penicillin-streptomycin | Gibco/Thermo Fisher | Cat#15140122 |
| HyClone SFX insect cell culture media | Cytiva/Fisher Scientific | Cat#SH3027802 |
| TNM-FH insect cell media | Sigma-Aldrich | Cat#T1032 |
| Trypsin treated with N-tosyl-L-phenylalanine chloromethyl ketone (TPCK) | Sigma-Aldrich | Cat#4370285 |
| L-glutamine | Gibco/Thermo Fisher | Cat#25030149 |
| **Critical commercial assays** |        |            |
| SigmaFast OPD Peroxidase Substrate | Sigma-Aldrich | Cat#P9187 |
| pFastBac Dual Expression Vector | Gibco/ThermoFisher | Cat#10712024 |
| **Experimental models: Cell lines** |        |            |
| Madin-Darby Canine Kidney (MDCK) cells | American Type Culture Collection (ATCC, VA, USA) | MDCK.2, Cat#CRL-2936 |
| Trichoplusia ni High Five (BTI-TNSB1-4) insect cells | American Type Culture Collection (ATCC, VA, USA) | Cat#CRL-10859 |
| Spodoptera frugiperda Sf9 insect cells | American Type Culture Collection (ATCC, VA, USA) | Cat#CRL-1711 |
| **Experimental models: Organisms/strains** |        |            |
| Specific-pathogen-free (SPF) chicken eggs | Canadian Food Inspection Agency (CFIA) | N/A |

(Continued on next page)
RESOURCE AVAILABILITY

Lead contact
Inquiries about any additional information pertaining to the study and requests for resources and reagents should be directed to the lead contact, Matthew S. Miller (mmiller@mcmaster.ca).

Materials availability
This study did not generate new unique reagents.

Data and code availability
- All raw data used to perform analyses in this article are available in tables found in the supplementary information.
- No original code was generated.
- Any additional information required to reanalyse the data reported in this paper is available from the lead contact upon request.

EXPERIMENTAL MODEL AND SUBJECT DETAILS

Study setting and participant recruitment
This study was nested within the multi-centre cluster randomized controlled trials (cRCT) of annual influenza vaccination in Hutterite communities. Selection of samples from the cRCT was done randomly, ensuring that the comparison groups were age- and gender-matched (see Table S1). Specifically, we included participants with a complete history of repeated vaccinations, who did not receive any external influenza vaccines and never tested PCR-positive for influenza A or B within the study period. Prior to initiation of this study, these communities had very low rates of vaccine uptake. The study also preceded universal recommendation of seasonal influenza vaccination for children in Canada. Therefore, the participants were unlikely to have received an influenza vaccine prior to study enrollment. For analysis of repeated vaccination with IIV (Figure 1A), we used annually collected pre-vaccination samples from the 2008-09, 2009-10 and 2012-13 seasons. The control group for this study included individuals who either received a hepatitis A vaccine, or no vaccine. No differences in immune responses were detected between the hepatitis A-vaccinated participants and the unvaccinated participants, and so they were grouped together for the purposes of this study. To assess the impact of vaccine formulation (IIV vs. LAIV, Figure 1B) we used samples collected at pre-vaccination day 0 and post-vaccination day 28 during the 2014-15 season. Vaccines were administered annually in October; participants’ records were screened for a period of 3 months post-vaccination to allow for reporting of any external vaccination and for reporting of PCR-confirmed influenza. The IIV and LAIV vaccinees received a 0.5-mL intramuscular injection of Vaxigrip (Sanofi Pasteur) or a 0.2-mL dose of intranasal FluMist (MedImmune), respectively. The following influenza strains were used in the standard-of-care trivalent vaccines administered over the study period: A/Brisbane/59/2007(H1N1), A/Brisbane/10/2007 (H3N2), B/Florida/4/2006-like virus (2008-09); A/Brisbane/59/2007 (H1N1), A/Brisbane/10/2007 (H3N2) and B/Brisbane/60/2008 (2009-10); A/California/7/2009 (H1N1), A/Perth/16/2009 (H3N2), B/Brisbane/60/2008 (2010-11) and A/California/7/2009 (H1N1)pdm09, A/Texas/50/2012 (H3N2), B/Massachusetts/2/2012 (2014-15) (Table 1). The control group consisted of participants who received either hepatitis A vaccine (N = 24) or no vaccine (N = 7) during the study period (Figure 1A); no differences in any of the assessed demographic or immune parameters were seen between these two subsets within the control group. All study procedures were approved by the McMaster University Research Ethics Review Board.

Cell lines
Madin-Darby Canine Kidney (MDCK) cells (ATCC, VA, USA) were maintained in Dulbecco Modified Eagle Medium (DMEM, Gibco) supplemented with 10% heat-inactivated fetal bovine serum (FBS, Gibco) and 100 U/mL penicillin-streptomycin (Gibco) at 37 °C.
in 5% CO2. *Trichoplusia ni* High Five (ATCC) insect cells were maintained in HyClone SFX insect cell culture media (Cytiva/Fisher Scientific) in shaker flasks shaking at 70 rpm at 28 °C. Spodoptera frugiperda Sf9 insect cells (ATCC) were maintained as adherent cultures in Roux flasks in full TNM-FH insect cell media (Sigma) supplemented with 10% FBS, 1% penicillin-streptomycin and 0.1% Pluronic F68 solution at 27 °C.

**Cal/09 H1N1 and chH5/1 N3 viruses**

A/California/04/2009 (Cal/09) H1N1 virus was kindly gifted by Dr. Peter Palese (Icahn School of Medicine at Mount Sinai, NY). The chH5/1 N3 virus was engineered to express ch5/1, a fusion of the A/Vietnam/1203/04 H5 head domain with PR8 group 1 stalk and neuraminidase N3. The viruses were propagated in 10-day-old embryonated specific-pathogen-free (SPF) chicken eggs sourced from Canadian Food Inspection Agency (CFIA) and maintained at 37°C in a dry incubator.

**Recombinant viral antigens**

All viral antigens were made using the baculovirus expression system and constructs kindly gifted by Dr. Peter Palese and Dr. Florian Krammer (Icahn School of Medicine at Mount Sinai, NY). The chH6/1 HA construct was composed of a A/Mallard/Sweden/81/02 H6 head domain and a A/Puerto Rico/8/1934 (PR8) group 1 stalk. Briefly, construct sequences were cloned into pFastBac Dual Expression Vector (Gibco/ThermoFisher) plasmids containing a C-terminal trimerization domain and hexahistidine tag, followed by transformation into DH10Bac bacteria (Gibco/ThermoFisher). The resulting bacmids were transfected into Sf9 cells to generate recombinant baculovirus, which was subsequently used to infect High Five insect cells. Viral supernatants were harvested via centrifugation at 5500g for 20 min at 4 °C, and proteins purified by Ni-resin affinity chromatography.

**METHOD DETAILS**

**Sample collection and processing**

Participant blood (5 mL) was collected into gold top serum separator tubes (BD) by venipuncture and centrifuged at 1300 × g for 15 min to extract serum. Nasal flocked swabs were collected into universal transport media (COPAN Diagnostics). All samples were stored at −80°C prior to analyses. Serum was inactivated by either heating at 56°C for 30 min prior to ELISA, or by a combination of 56°C and trypsin-heat-periodate treatment prior to MNT and HAI assays.

**Microneutralization (MNT) assays**

Participant sera were serially diluted 2-fold to create a 40 to 5120-fold dilution series using MEM supplemented with 1x penicillin-streptomycin, 25 nM HEPES, and TPCK-treated trypsin (0.25 - 1 μg/mL) (Sigma-Aldrich). The diluted sera were then incubated with the chH5/1 N3 virus at a concentration of 400 PFU/well at RT for 30 min. The virus-serum mix was then added to MDCK cells plated in a 96-well plate at ~80% confluency and incubated at 5% CO2 for 1 h at 37°C. The plate was then washed and supplemented with DMEM containing 0.2 M L-glutamine, 100U/mL penicillin and 1 μg/mL streptomycin overnight at 5% CO2, 37°C, followed by fixation with 80% acetone. Influenza virus nucleoprotein (NP) was measured using biotin-conjugated anti-NP antibody (EMB Millipore). SigmaFast OPD (Sigma-Aldrich) was used as HRP substrate and luminescence was measured using the SpectraMax i3 plate reader at OD490. A MNT endpoint titer, serum dilution resulting in at least 50% inhibition of infectivity, was defined as the reciprocal of the lowest serum dilution factor, for which normalized signal remained below the threshold of detection; endpoints outside the assay’s lower limit of detection (<40) were assigned an endpoint titer of 20.

**Haemagglutination inhibition (HAI) assays**

Serum was mixed with the Cal/09 H1N1 virus and incubated for 30 min at RT. Chicken erythrocytes in Alsever’s solution (sourced from CFIA) were then added to the plate and incubated at 4°C for 45 min. An HAI titer was defined as the highest serum dilution factor leading to HAI activity; samples without any detectable HAI activity were assigned a titer of “5”.

**Influenza virus antibody ELISA**

96-well plates were coated with 2 μg/mL of Cal/09 or chH6/1 HA, followed by 4°C incubation overnight and blocking with 5% non-fat milk in PBS-T. Sera were added to the plate in a 2-fold serial dilution series ranging from 100- to 6400-fold. Mucosal samples were assayed undiluted. Plates were then incubated for 1 h at RT, HRP-conjugated goat anti-human IgG or IgA were added. SigmaFast OPD (Sigma-Aldrich) was used as HRP substrate and luminescence was measured using the SpectraMax i3 plate reader at OD490. A detection threshold was defined as 3 standard deviations above the average background signal. Serum IgA and IgG endpoint titers were defined as the highest serum dilution factor for which normalized OD490 values remained above the background threshold; endpoints outside the assay’s limit of detection (i.e. <100 or >6400) were assigned endpoint titer values of 50 and 25600, respectively. Mucosal ELISA readouts were reported as normalized OD490 values; post-normalization values < 0 were substituted with assay-specific lowest limit of detection values for the geometric mean fold change (GMFC) and log2 calculations.
Statistical analyses were performed using IBM SPSS v.27 and GraphPad Prism v.7. Two-sided Mann-Whitney U and Fisher’s exact tests were used to assess demographic differences among the participant groups in Tables S1 and S4. Data were scaled via log₂ transformation. Geometric means and 95% confidence intervals were calculated for each time point in Figures 2, 3, 4, 5, S2, and S3. GMFC were calculated as ratios of geometric means (i.e. pre-/post-vaccination) for each parameter. Paired analyses were done using the paired t-test in Figures 2, 3, 4, 5, S1, S2, and S3. Differences between the LAIV and IIV groups were assessed using the unpaired t test in Figures 2, 3, 4, 5, and S2. Correlation analyses were performed using the Pearson correlation coefficient test and lines of best fit were derived via linear regression in Figures 2, 3, 4, 5, 6, and S2. Differences and correlations between variables were considered significant at $p < 0.05$. 