

Live Attenuated Influenza Vaccine

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Abstract Cold-adapted Ann Arbor based live attenuated influenza vaccine (LAIV) has been available in the USA since 2003. The vaccine is efficacious against influenza infection. Features of LAIV include: easy administration suitable for mass immunization, cross-reactivity to drifted strains for broader coverage, and establishment of herd immunity for control of influenza spread. Annual seasonal LAIV now contains four strains against influenza A H1N1, H3N2, influenza B-Victoria, and B-Yamagata lineages that are co-circulating in humans. LAIV played a significant role in protecting the public from the 2009 H1N1 pandemic and has been evaluated for pandemic preparedness. Pandemic vaccines including influenza H2, H5, H6, H7, and H9 subtypes have been produced and evaluated in preclinical and small-scale phase I clinical studies. This review summarizes the current status and perspectives of seasonal and pandemic LAIV.

Abbreviations

LAIV	Live attenuated influenza vaccine
T/LAIV	Trivalent live attenuated influenza vaccine
Q/LAIV	Quadrivalent live attenuated influenza vaccine
pLAIV	Pandemic live attenuated influenza vaccine
TIV	Trivalent inactivated influenza vaccine
IIV	Inactivated influenza vaccine
MDV-A	Master donor virus for influenza A vaccines
MDV-B	Master donor virus for influenza B vaccines
<i>ca</i>	Cold adapted
<i>ts</i>	Temperature sensitive
<i>att</i>	Attenuated

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<i>wt</i>	Wild-type
RG	Reverse genetics
PCKC	Primary chicken kidney cells
NT	Nasal turbinates
HAI	Hemagglutination inhibition
NW	Nasal wash

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1 Introduction

Influenza viruses cause significant morbidity and mortality, leading to more than 100,000 hospitalizations and 3,000–49,000 deaths annually (Thompson et al. 2003). Influenza viruses undergo constant genetic drift resulting in emergent antigenic variants that can escape immunity to HA and NA antigens of previously circulating strains. Therefore, influenza vaccines must be updated annually to match the contemporary strains in order to provide optimal protection. Trivalent LAIV (T/LAIV, H1N1, H3N2, one B strain) in a frozen formulation was approved by the US FDA in 2003 for healthy persons 5–49 years of age and the next generation of a liquid formulation was approved for 2–49 year olds in 2007. LAIV is the first new influenza vaccine, as well as the first nasally administered vaccine of any kind for human use, in the USA since the introduction of injectable trivalent influenza vaccine (TIV) in the 1940s (Grabenstein et al. 2006). The formulation of

the vaccine was recently changed from trivalent to quadrivalent to account for the epidemiology of B strains in circulation. Two antigenically distinct lineages of influenza B viruses have circulated globally since 1985 and vaccines against one lineage do not offer cross-protection against the other lineage (Rota et al. 1990). In order to provide broader coverage of influenza B viruses an additional B strain is now included in the vaccine. T/LAIV was therefore discontinued in 2013 with the approval and marketing of a quadrivalent LAIV (Q/LAIV, H1N1, H3N2, two B strains).

The development of LAIV is the culmination of over 40 years of collaborative research and development between University of Michigan and scientists from the National Institutes of Health (NIH) and the biopharmaceutical industry (Wyeth, Aviron, and MedImmune). Following the approval in the USA in 2003, T/LAIV was approved in Israel, South Korea, United Arab Emirates, Mexico, and Macau for individuals 2–49 years of age, in Canada for individuals 2–59 years of age and in the European Union for individuals 2–17 years of age. In June 2014, the Advisory Committee on Immunization Practices (ACIP) made a preferential recommendation of LAIV for healthy children ages 2 to 8 years old in the USA.

2 Development of Cold-Adapted Ann Arbor Donor Viruses

LAIV is developed based on two cold-adapted master donor viruses, A/Ann Arbor/6/60 (H2N2) for influenza A vaccines (MDV-A) and B/Ann Arbor/1/66 for influenza B vaccines (MDV-B). Each donor virus donates the cold adapted (*ca*), temperature sensitive (*ts*), and attenuated (*att*) phenotype to the 6:2 reassortant vaccine viruses that contain six internal protein gene segments of MDV and the HA and NA surface proteins of the wild-type (*wt*) influenza virus. Both MDV-A and MDV-B were developed by Dr. John Maassab at the University of Michigan in the 1960s (Maassab 1967). Influenza A/Ann Arbor/6/60 was subjected to serial in vitro passage at gradually reduced temperature in primary chicken kidney cells (PCKC), 2 passages (2x) at 36–37 °C, 7x at 33 °C, 7x at 30 °C, 7x at 25 °C, 6x plaquing at 25 °C and 3x amplification in embryonated chicken eggs. Influenza B/Ann Arbor/1/66 was passaged less extensively, two passages (2x) at 36–37 °C, 2x at 33 °C, 5x at 27 °C, 6x at 25 °C, 7x plaquing at 25 °C, and 3x amplification in eggs. During cold passage, the MDV-A and MDV-B acquired a number of genetic changes in multiple gene segments, which confer the *ca*, *ts*, and *att* phenotypes that can be imparted to 6:2 reassortant vaccine strains. The *ca* phenotype reflects efficient viral replication at a lower temperature of 25 °C, while most *wt* influenza viruses do not replicate well at this temperature. The *ts* phenotype reflects restricted replication at a higher temperature at which most *wt* viruses can replicate well. The shut-off temperature of MDV-A and MDV-B is different: 39 °C for MDV-A and 37 °C for MDV-B. The *att* phenotype can be measured in the ferret model, viral replication is detected in the nasal tissues but not in the lungs of

ferrets intranasally infected with the vaccine virus. The *calts/att* phenotypes provide safety features of the reassortant vaccine strains that can replicate in the cooler nasal tissues but not at the higher core body temperature of the lungs.

3 Genetic Basis of *calts/att* Phenotypes of the Vaccine Donor Viruses

3.1 MDV-A

By comparing viral genomic sequences between the *wt* and *ca* A/Ann Arbor/6/60 strains and introduction of each of the mutations individually and combination, the amino acids that confer the *ts* phenotype have been precisely mapped to three residues (E391, G581, T661) in PB1, one residue (S265) in PB2 and one residue in NP (G34) proteins (Table 1) (Jin et al. 2003). These five loci in the PB1, PB2, and NP genes also confer the *att* phenotype (Jin et al. 2004). The *ca* phenotype of MDV-A could not be experimentally mapped as the available *wt* A/Ann Arbor/6/60 strain had been passaged extensively in tissue culture and also grows well at 25 °C. Since the *ts* and *att* phenotypes are specified by five residues in three gene segments, the chance for the vaccine virus to revert to *wt* phenotype is extremely low, explaining genetic stability of the vaccine strains.

Limited studies have been conducted to understand the molecular mechanism of these loci in specifying viral phenotypes. By minigenome analysis, the five loci have been shown to greatly reduce viral RNA-dependent polymerase activity of AA *ca* at the nonpermissive temperature of 39 °C (Jin et al. 2004). During viral infection *in vitro*, the vaccine virus can initiate single cycle replication, but multicycle viral replication at 39 °C is significantly reduced. vRNA synthesis and translocation of viral RNP from nucleus to cytoplasm are reduced. In addition, incorporation of the M1 protein into virions is significantly reduced, resulting in irregular viral morphology (Chan et al. 2008).

3.2 MDV-B

The *calts/att* phenotypes of MDV-B have been mapped by reverse genetics (RG) (Chen et al. 2008; Hoffmann et al. 2005). The *ts* loci are specified by three amino acids in the PA and NP proteins: M431 in the PA, A114, and H410 in the NP (Table 1). These three residues and Q159 and V183 residues in the M gene segment contribute to the *att* phenotype. Five residues in three segments confer the *ca* phenotype: R630 in PB2, M431 in the PA, A114, H410, and T509 in the NP. A total of seven loci distributed in four gene segments of MDV-B control the *calts/att* phenotypes, making the vaccine donor genetically stable. These loci not only reduce viral polymerase function but also affect virus assembly and release at the restricted temperature (Chan et al. 2008).

Table 1 Genetic loci of MDV-A and MDV-B

Segment	MDV-A		MDV-B	
	Amino acid	Phenotype	Amino acid	Phenotype
PB2	S265	<i>ts/att</i>	R630	<i>ca</i>
PB1	E391 G581 T661	<i>ts/att</i>	–	–
PA	–	–	M431	<i>ca/ts/att</i>
NP	G34	<i>ts/att</i>	A114 H410 T509	<i>ca/ts/att</i> <i>ca/ts/att</i> <i>ca</i>
M	–	–	N159 V183	<i>att</i>
NS	–	–	–	–

–: No role in the *ca/ts/att* phenotypes

4 LAIV by Reverse Genetics and Yearly LAIV Production Process

LAIV is manufactured in specific pathogen free (SPF) embryonated chicken eggs. Human influenza viruses normally do not replicate well in eggs and viral replication in eggs often results in mutations in the HA receptor binding region. The quasi-species of HA sequence variations from egg adaptation have different abundance and biological impact. While certain changes improve viral growth in eggs without affecting viral antigenicity and immunogenicity, other changes are not suitable for a vaccine. It is therefore critical to evaluate multiple candidate vaccines in order to select an appropriate HA variant for vaccine production. The application of reverse genetics (RG) technology to the production of 6:2 reassortant vaccine viruses has revolutionized the vaccine seed production process. Traditional classical reassortment, which is a very time consuming and unpredictable method, has been replaced by the use of RG for LAIV production since the 2008–2009 influenza season. The 6:2 reassortant vaccine virus contains six internal protein gene segments from MDV-A or MDV-B and the surface HA and NA glycoprotein gene segments of a *wt* virus (Fig. 1).

During vaccine seed production, each candidate vaccine variant is evaluated for yield in eggs, antigenicity is assessed by reactivity with a reference antiserum from ferrets immunized with *wt* virus, and immunogenicity is assessed by examining serum antibody levels achieved in ferrets following intranasal administration of vaccine virus variants. Table 2 summarizes the amino acid sequence changes that have been frequently detected in the HA and their impact on viral biological activities (for a recent review, please see Jin and Chen 2014). Most of the changes do not affect viral antigenicity and immunogenicity, however, several changes

Fig. 1 Generation of reassortant LAIV by RG. The six plasmids of the internal protein gene segments of MDV and two plasmids encoding *wt* HA and NA gene segments are electroporated into Vero cells. Virus recovered from the transfected cells is amplified in embryonated chicken eggs

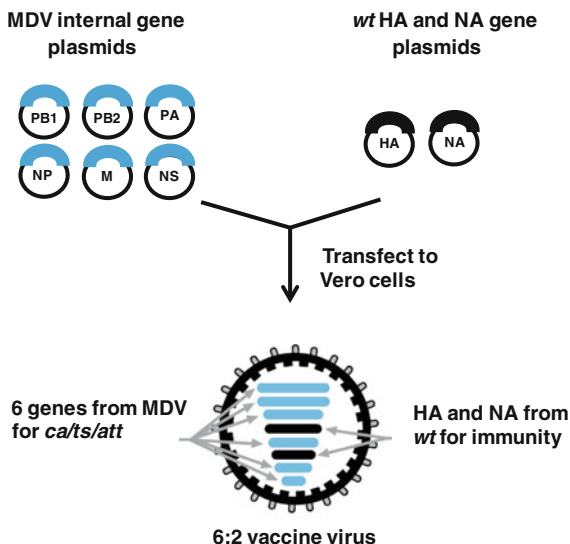


Table 2 Common egg adaptation sites in the HA of egg-grown influenza viruses

Influenza viruses	HA amino acid ^a	Comments related to residues in bold
H1N1	119, 125, 127, 153–155 , 186, 209, 222, 223	153–155 residues alter viral antigenicity
H3N2	138, 156 , 183, 186, 189, 190, 193, 194 , 195, 196, 219, 226	156 is a major antigenic site 194 affects viral replication in ferrets
B-Vic	197, 199	Glycosylation site
B-Yam	196, 198	Glycosylation site

^a Amino acid numbering based on HA sequence of each subtype

identified in the HA have been shown to reduce viral antigenicity or to render vaccine virus less immunogenic (Chen et al. 2010a, b).

Once a LAIV seed is selected, the vaccine virus is purified by one round of limited dilution in eggs to produce a Master Virus Seed (MVS) that is used for bulk production of monovalent vaccine. Each vaccine lot is subjected to a lot release test to ensure that no adventitious agents are present. The four vaccine viruses, H1N1, H3N2, B-Yamagata, and B-Victoria, are formulated to contain $10^{7.0 \pm 0.5}$ FFU (fluorescent focus units)/strain in 0.2-mL in a nasal sprayer. If a new vaccine virus component is incorporated into the annual seasonal vaccine, the vaccine is tested in a small-scale safety trial prior to final release of the vaccine for widespread use.

5 Preclinical Studies of LAIV

Extensive studies have been performed to evaluate vaccine viruses for their attenuated replication, immunogenicity, and protection against *wt* influenza virus challenge infection in the ferret model. Vaccine viruses have been shown to be able to replicate in the nasal turbinates (NT) but not in the lungs of vaccinated ferrets. Although vaccine strains in general elicit hemagglutination inhibition (HAI) antibodies at levels lower than that induced by homologous *wt* virus (MedImmune data), they provide significant protection against replication of homologous challenge virus in the upper and lower respiratory tract (LRT) of ferrets.

Ferret studies have been conducted to compare T/LAIV with Q/LAIV in protection against *wt* challenge infection to address vaccine interference (Bandell et al. 2011). Q/LAIV is comparable to T/LAIV in vaccine-induced protective immune responses against *wt* virus replication in the upper and LRT of vaccinated ferrets (Fig. 2). The vaccine strains A/California/7/2009 (A/CA/09, H1N1), A/Perth/16/2009 (A/Perth/09, H3N2), B/Brisbane/60/2008 (B/Bris/08, Victoria lineage), and B/Wisconsin/1/2010 (B/Wis/10, B/Yamagata lineage) had comparable immunogenicity given in either Q/LAIV or T/LAIV formulation, elicited robust antibody responses to each vaccine component and fully protected ferrets from *wt* virus challenge in the lungs after two doses of vaccine. Each vaccine strain also offered significant protection from *wt* virus replication in the NT. These studies indicate that LAIV strains in multivalent vaccine formulations do not demonstrate evidence of viral or immune interference affecting efficacy of each vaccine component.

6 Clinical Studies of LAIV

Clinical studies of Ann Arbor (AA) *ca*-based LAIV strains in the USA include monovalent, bivalent including two type A strains, trivalent including H1N1, H3N2 and a B strain, and quadrivalent including two type A and two B strains (for review see Murphy and Coelingh 2002). T/LAIV has been evaluated in more than 73 clinical research trials completed worldwide in >141,000 people ranging in age from 6 weeks to >90 years. Approximately 80 million doses have been distributed for commercial use since the initial US licensure in 2003 up through March 2014 (LAIV Scientific Product Monograph 2013–2014, MedImmune, Gaithersburg, MD).

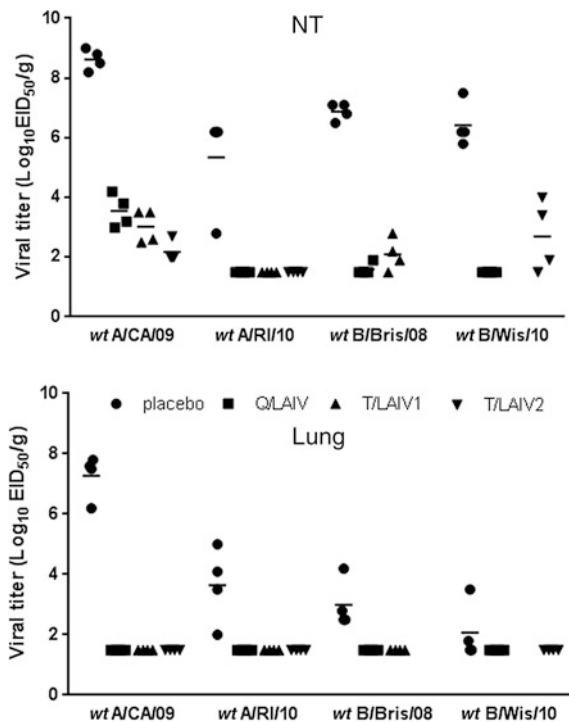


Fig. 2 Challenge virus titers in NT and lung of ferrets vaccinated with Q/LAIV or T/LAIV. Ferrets were vaccinated with sucrose phosphate (SP) buffer, T/LAIV1 (A/CA/09, A/Perth/09, B/Bris/08), T/LAIV2 (A/CA/09, A/Perth/09, B/Wis/10), or Q/LAIV (A/CA/09, A/Perth/09, B/Bris/08 and B/Wis/10) in two doses one month apart. The ferrets were challenged with indicated homologous *wt* viruses with the exception of the H3N2 A/Rhode Island/2010 strain as *wt* A/Perth/09 did not replicate well in ferret lungs. Viral titers in the NT and lung tissues were determined by 50 % egg infectious dose (EID₅₀) assay. The limit of detection of the assay was 1.5 log₁₀EID₅₀/g

6.1 Safety

LAIV is generally well tolerated and safe (reviewed by Ambrose et al. 2011). The vaccine viruses infect and replicate in cells lining the nasopharynx of the recipient to induce immunity, but are not able to replicate in the LRT due to their *ts* and *att* phenotypes. The most common solicited adverse reactions are runny nose or nasal congestion (ages 2 to 49 years), fever >100 °F (ages 2–6 years), and sore throat (18–49 years). The rate of headache and tiredness in LAIV recipients is higher than in a placebo control group but is similar to TIV recipients (Baxter et al. 2012a, b; Toback et al. 2013). One study showed that LAIV was associated with an increased rate of all-cause hospitalization among children aged 6–11 months and an increased rate of medically attended wheezing in children aged 6–23 months (Belshe et al. 2007). For this reason, LAIV is not approved for children younger than 24 months of age.

6.2 Transmission and Genetic Stability

Vaccine viruses can be cultured from nasal secretions in the first few days after vaccination. The relationship of viral replication in a vaccine recipient to transmission of vaccine viruses to other individuals has not been well established. LAIV is poorly transmissible because most infection is not symptomatic and this decreases the likelihood of viral spread via cough or sneezing. In addition, levels of vaccine virus replication in nasal tissues are much lower than *wt* virus even in seronegative children making its spread to contacts very inefficiently (Murphy and Coelingh 2002). In studies performed to date, viruses shed from vaccine recipients have been consistently phenotypically and genotypically stable, maintaining the *ca*, *ts*, and *att* phenotypes (Cha et al. 2000; Vesikari et al. 2006). LAIV has been shown to be poorly transmissible to spouses, roommates, and household members under a variety of circumstances in small clinical trials (Murphy and Coelingh 2002). Eighty percent of trivalent LAIV recipients who were 6–36 month old children in a day care setting, shed at least one vaccine strain, with a mean duration of shedding of 7.6 days ranging from 1 to 21 days (Vesikari et al. 2006). Transmission of vaccine viruses from vaccine recipients to placebo subjects was a rare event. The *ca* and *ts* phenotypes were preserved in all recovered viruses tested ($n = 135$ tested of 250 strains isolated at the local laboratory). The probability of a young child acquiring vaccine virus after close contact with a single trivalent LAIV vaccinee in a day care setting was 0.58 % (95 % CI: 0, 1.7) based on the Reed Frost model (Longini et al. 1982). With documented transmission of type B virus in one placebo subject and possible transmission of type A virus in four placebo subjects, the maximum probability of acquiring a transmitted vaccine virus was estimated to be 2.4 %.

6.3 Efficacy

The efficacy and effectiveness of an influenza vaccine can be evaluated by three criteria: (1) comparison of culture-positive influenza infection rates, which are most feasible in young children because they readily shed influenza virus (i.e., vaccine efficacy); (2) a 4-fold antibody increase from baseline levels, which is subject to inherent bias from prior vaccine or natural disease exposure, and therefore is a method of limited value; or (3) observations of clinical events or “medically attended acute respiratory illness” (MAARI) which is a less specific endpoint than culture-confirmed influenza illness, and results in effectiveness point estimates that are significantly lower than efficacy estimates. Adults shed virus in low quantity and for a short duration and thus clinical trials in adults are more commonly conducted using clinical endpoints (Belshe et al. 2004). LAIV efficacy trials in the pediatric population consist of nine controlled studies comprising over 20,000 infants and toddlers, children, and adolescents, during seven influenza

seasons (summarized in LAIV Scientific Product Monograph 2013–2014, Med-Immune, Gaithersburg, MD). Four placebo-controlled studies were conducted to include revaccination in a second season. Overall, LAIV has an efficacy of 62–93 % against antigenically matched strains and 49–93 % against all strains in children 15–71 months of age. LAIV has demonstrated superiority compared to TIV in 3 active-controlled studies in children. LAIV is about 44 % (range 34.7–52.7 %) better than TIV for matched strains and 31.9–54.9 % better than TIV for all strains. In one of the largest field efficacy trials (MI-CP111), LAIV was shown to be more efficacious than inactivated TIV in children 6–59 months of age (Belshe et al. 2007). Because of the higher efficacy of LAIV in children, LAIV is preferentially recommended in the UK, Germany, Israel, Canada, and Sweden for children of various age groups (Ambrose et al. 2012), and recently for 2–8 year old children in the USA.

LAIV efficacy in adults has been demonstrated in two efficacy trials (Nichol et al. 1999b; Treanor et al. 2000). In the first trial, LAIV was shown to reduce laboratory-documented influenza illness by 85 % compared to TIV in a challenge study conducted in healthy adults 18–41 years of age who were presumed to be susceptible to at least one strain included in the vaccine based on prevaccination antibody titers (Treanor et al. 2000). In the second, larger trial, LAIV recipients exhibited significant reduction in days of febrile illness, missed work, health care provider visits, and antibiotic usage. The efficacy of LAIV and TIV can be affected by a number of factors, such as the age and health of the vaccine recipients and the extent of antigenic similarity between the vaccine strains and circulating strains. In a study conducted by Monto et al. (Monto et al. 2009), LAIV was 50 % less efficacious than TIV in reduction of laboratory-confirmed influenza during the 2008–2009 influenza season when an H3N2 virus was the predominant circulating strain. Based on a subgroup analysis of subjects 50–64 years of age in the study by Nichol et al. (1999a), LAIV was not approved for this age group in the USA. A later study showed that LAIV offered statistically significant protection against culture-confirmed influenza in adults ≥ 60 years of age (De Villiers et al. 2009).

6.4 Immunogenicity

The immunogenicity of 19 different LAIV strains was studied over a period of 25 years at various locations and in different populations (Murphy and Coelingh 2002), and annual commercial vaccines have been evaluated over the past 10 years. Protection against influenza generally correlates with serum IgG hemagglutination-inhibiting antibodies (HAI), especially in seronegative children. After two doses of LAIV, children who were presumed to be susceptible to at least one strain included in the vaccine based on prevaccination antibody titers, mounted an adequate HAI response (>90 % seroconverted to type A/H3 and B strains, and 60–90 % to type A/H1 strain) (Belshe et al. 1998, 2000; Zangwill 2003). Antibodies persisted for 5–8 months after vaccination with LAIV, and

protection generally persisted for at least 1 year (Zangwill 2003). In a study of young children, protective efficacy lasted for the duration of the influenza season and as late as 5.5–13 months after the second dose (Ambrose et al. 2008; Tam et al. 2007). In adults, the serologic response has been less robust (<35 % for A/H3 and B and 60–90 % for A/H1), and the correlates of immunity may be related to other immune responses (Gorse et al. 1995; Tomoda et al. 1995; Zangwill 2003). LAIV may be more effective than IIV in inducing a nasal IgA response that is important for viral clearance and recovery, whereas IIV vaccine more consistently elicits serum HA antibodies in adults (Beyer et al. 2002; Cox et al. 2004; Renegar et al. 2004). The role of cell-mediated immune responses in the protection of young children against influenza was studied in a large randomized, double-blind, placebo-controlled dose-ranging efficacy trial with 2,172 children of 6 to <36 months old in Philippines and Thailand (Forrest et al. 2008). LAIV was found to elicit substantial CMI responses as measured by interferon-gamma ELISPOT assay that correlated with protection. Another study conducted in children showed that LAIV induced cell-mediated responses including CD4(+), CD8(+), and $\gamma\delta$ T cells that are relevant for broadly protective heterosubtypic immunity (Hoft et al. 2011). In a study conducted in young adults, although TIV induced higher levels of vaccine-specific plasmablasts and plasmablast-derived polyclonal antibodies (PPAb) than LAIV, LAIV induced a greater vaccine-specific IgA plasmablast response as well as a greater plasmablast response to the conserved influenza nucleoprotein and better cross-reactivity to heterologous strains than TIV (Sasaki et al. 2014).

7 Pandemic LAIV

There are 18 known HA and 11 known NA subtypes of influenza A viruses in nature; 16 HA and 9 NA subtypes have been isolated from waterfowl and shorebirds, and a variety of subtypes have been isolated from other animal species including pigs, horses, and dogs. Animal influenza viruses are the source from which novel HA and NA subtypes are introduced into the human population, by reassortment with other animal or human influenza viruses or direct infection of humans. In the last century, influenza pandemics occurred in 1918, 1957, and 1968 and the first pandemic of this century occurred in 2009. Each of these pandemics was associated with significant morbidity and mortality. Since 1997, animal H5N1, H6N1, H7N7, H7N3, H7N9, H9N2, and H10N8 viruses have caused human infections but have not spread efficiently from person to person. The pandemics and the sporadic emergence of animal viruses into humans underline the need for the generation of pandemic influenza vaccines and their evaluation in humans. The criteria for licensure of currently licensed inactivated influenza vaccines are protective antibodies directed primarily against the HA, the major protective antigen of the virus that induces neutralizing antibody and/or demonstrated efficacy.

LAIV have several attributes related to safety, immunogenicity, cross-protection against antigenic drift strains, high yield, and needle-free administration that make them attractive candidates for control of pandemic influenza. LAIV generally induce broadly cross-reactive protection (Coelingh et al. 2014; Murphy and Coelingh 2002), which may be a useful feature in the event of a pandemic if a vaccine generated from the emerging pandemic strain is not immediately available. Importantly, the infrastructure for manufacture and distribution of a LAIV exists. Therefore, the United States NIH and MedImmune undertook a joint effort to develop and evaluate LAIV bearing the HA and NA genes from animal influenza viruses on the MDV-A backbone as pandemic LAIV (pLAIV) candidates. Our approach includes: (1) generation of a pLAIV bearing an HA and appropriate NA from an animal influenza virus on the attenuated MDV-A background; (2) evaluation of the attenuation, immunogenicity, and protective efficacy of the candidate vaccine in animal models; (3) preparation and qualification of a clinical lot of each pandemic vaccine candidate; (4) evaluation of the safety, infectivity, and immunogenicity of each candidate in humans; (5) storage of human sera obtained from vaccinees to determine cross-reactivity with the newly emerged pandemic viruses; and (6) storage of seed viruses for use in the manufacture of vaccine to prevent disease caused by related pandemic viruses that do emerge such that vaccine manufacture can be initiated with pretested vaccines without delay.

A theoretical concern associated with the use of a pLAIV bearing genes derived from an animal influenza virus is the risk of reassortment of the vaccine virus with a circulating influenza virus, resulting in a novel subtype of influenza that could spread in the human population. Although such a reassortment event may not be of great significance in the face of widespread disease caused by a pandemic influenza strain, it would be an unfavorable outcome if the threatened pandemic did not materialize. This risk would be carefully considered by public health authorities before a decision is made to introduce a live attenuated vaccine in a threatened pandemic. With the exception of one H9N2 *ca* virus, the pLAIV viruses were generated by RG. The HAs of highly pathogenic avian influenza (HPAI) H5 and H7 viruses were modified to remove the multibasic amino acid cleavage motif that is a known virulence determinant for poultry.

7.1 Preclinical Studies

We have developed pLAIV candidates against 6 different subtypes (H1, H2, H5, H6, H7, and H9) (Chen et al. 2003, 2009a, b, 2011b, 2014; Joseph et al. 2008; Min et al. 2010; Suguitan et al. 2006). The genetic loci responsible for the *ts* and *att* phenotypes associated with the MDV-A were confirmed in each of the pLAIV viruses. The replication of the pLAIV viruses was evaluated in the respiratory tract of ferrets 3 days following intranasal administration of 10^7 TCID₅₀ of viruses. As discussed previously, highly restricted replication in the LRT of ferrets defines the *att* phenotype of the pLAIV strains (Table 3). Some pLAIV viruses (H2, H5, and H7N3)

Table 3 Replication of intranasally administered pLAIV in the respiratory tract of ferrets and mice

Subtype/Vaccine virus	Virus titer (\log_{10} TCID ₅₀ /g) in the respiratory tract of indicated species			
	Ferrets ^a		Mice (peak titer) ^b	
	URT	LRT	URT	LRT
pH1N1 CA/7/2009 (Chen et al. 2011b)	3.4	1.7	2.0	1.8
H2N2 Ann Arbor/6/60 (Chen et al. 2009a)	5.9	$\leq 1.5^c$	5.0	4.3
H2N3 swine/MO/2006 (Chen et al. 2014)	5.0	$\leq 1.5^c$	4.3	3.1
H5N1 VN/1203/2004 (Suguitan et al. 2006)	4.1	$\leq 1.5^c$	2.6	4.1
H5N1 HK/213/2003	4.5	$\leq 1.5^c$	2.5	5.1
H6N1 teal/HK/W312/97 (Chen et al. 2009b)	2.1*	$\leq 1.5^c$	nd	nd
H7N3 BC/CN-6/2004 (Joseph et al. 2008)	4.7	$\leq 1.5^c$	4.4	2.4
H7N7 NL/219/2003 (Min et al. 2010)	2.8	$\leq 1.5^c$	3.3	3.3
H9N2 ck/HK/G9/1997 (Chen et al. 2003)	2.3	$\leq 1.5^c$	1.4**	$\leq 1.5^c$

^a Lightly anesthetized ferrets received intranasal administration of 10^7 TCID₅₀ (or fluorescent forming units) of virus and virus titers in the upper and lower respiratory tract were determined 3 days later

^b Lightly anesthetized mice received intranasal administration of 10^6 TCID₅₀ of virus and virus titers in the upper and lower respiratory tract were determined at serial time points 2, 3, and 4 days later. The peak titer achieved is indicated

^c Below the lower limit of detection

nd Not done, *pfu/g, **TCID₅₀/ml

replicated to high titer in the upper respiratory tract (URT) of ferrets, while others such as the H6N1, H7N7, and H9N2 vaccine viruses were restricted in replication in the URT. The pLAIV were also evaluated for replication in mice 2, 3, and 4 days following intranasal administration of 10^6 TCID₅₀ of virus (Table 3). The body temperature of mice is 37 °C, which is below the shut-off temperature of the ca viruses. Therefore, although the replication of the pLAIV viruses in mice is restricted compared to the corresponding *wt* viruses, they are not as restricted in replication in mice as they are in ferrets that have a core body temperature of ~39 °C.

The immunogenicity of the pLAIV was evaluated in mice and ferrets by measuring the serum antibody response by HAI and/or microneutralization (MN) assays against homologous and heterologous viruses following intranasal administration of one or two doses of pLAIV; neutralizing antibody (MN) titers are reported in Tables 4 and 5. The pH1N1 LAIV that was used for vaccine production incorporated two mutations in the HA gene that improved the yield in eggs and immunogenicity in ferrets (Chen et al. 2010a). The homologous antibody response to one dose of vaccine in mice and ferrets ranged from poor (VN04 H5N1 and H7N7 vaccines) to robust (H6N1 and H9N2 vaccines) (Tables 4 and 5) but a correlation between the magnitude of the antibody response and replication of the pLAIV in mice or ferrets was not apparent (Tables 3, 4 and 5). Notably, in all cases, a second dose of vaccine boosted serum antibody responses against homologous and heterologous viruses.

Table 4 Immunogenicity and efficacy of one or two doses of pLAIV in mice, evaluated against homologous and heterologous viruses of the same subtype

Subtype and vaccine virus	Immunogenicity ^a by MN (GMT)				Efficacy ^b against challenge infection									
	1 dose		2 doses		1 dose		2 doses		1 dose		2 doses		Heterologous ^c	
	Homologous	Heterologous	Homologous	Heterologous	URT	LRT	URT	LRT	URT	LRT	URT	LRT	URT	LRT
pH1N1 CA/7/2009 (Chen et al. 2011b)	53	nd	595	nd	4+	2+	nd	nd	5+	5+	5+	5+	nd	nd
H2N2 Ann Arbor/6/60 (Chen et al. 2009a, 2014)	nd	nd	80	14–17	5+	5+	2+ to 3+	2+ to 3+	5+	5+	5+	5+	nd	nd
H2N3 swine/MO/2006 (Chen et al. 2014)	57	10–19	2,874	55–94.1	5+	4+	5+	3+ to 5+	5+	5+	5+	5+	5+	5+
H5N1 VN/1203/2004 (Suguitan et al. 2006)	14	11–15	388	160–528	2+	2+	+ to 5+	+ to 5+	5+	5+	5+	5+	5+	5+
H5N1 HK/213/2003	59	10	1,056	19–61	5+	5+	3+ to 5+	2+	5+	5+	5+	5+	5+	5+
H6N1 teal/HK/W/312/97 (Chen et al. 2009b)	94	10–86	543	15–388	5+	5+	5+	5+	5+	5+	5+	5+	5+	5+
H7N3 BC CN-6/2004 (Joseph et al. 2008)	87	<20	470	39–60	5+	5+	2+ to 5+	3+ to 5+	5+	5+	5+	5+	5+	5+
H7N7 NLJ/219/2003 (Min et al. 2010)	14	21–24	191	153–285	2+	+ to 3+	3+ to 5+	+ to 3+	2+	+ to 3+	4+	5+	5+	5+
H9N2 ck/HK/G/9/1997 (Chen et al. 2003)	320 ^d	340 ^d	nd	nd	5+	5+	5+	5+	5+	5+	nd	nd	nd	nd

^a Serum neutralizing antibody titer measured in a microneutralization (MN) assay

^b Efficacy was assessed by reduction of replication of wild-type challenge virus compared to mock-vaccinated animals. — <10-fold reduction, + 10 to 99-fold reduction, 2+ 100 to 999-fold reduction, 3+ 1000 to 9999-fold reduction, 4+ >10,000-fold reduction, 5+ complete protection from replication

^c Heterologous challenge viruses for H2 vaccines: Japan/57 (H2N2), swine/MO/06 (H2N3); for H5 vaccines: HK/491/97, HK/213/2003; Vietnam/1203/2004, Indonesia/5/2005; for H6 vaccines: dk/HK/77, mallard/Alberta/85; for H7 vaccines: turkey/V/455/02 (H7N2), turkey/UT/24721-10/95 (H7N3), chicken/BC/04 (H7N3), Netherlands/219/03 (H7N7), tk/Italy/99 (H7N1), turkey/England/63 (H7N3); for H9 vaccines: HK/1073/99

^d HAI titer

nd/ Not done

Table 5 Immunogenicity and efficacy of pLAIV in ferrets, evaluated against homologous and heterologous viruses of the same subtype

Subtype and vaccine virus ^a	Immunogenicity ^b by MN (GMT)			Efficacy ^c against challenge infection						
	1 dose		2 doses		1 dose		2 doses		Heterologous ^d	
	Homologous	Heterologous	Homologous	Heterologous	URT	LRT	URT	LRT	URT	LRT
pH1N1 CA/7/2009 (Chen et al. 2011b)	854	nd	2,428	nd	5+	5+	nd	5+	5+	nd
H2N2 Ann Arbor/6/60 (Chen et al. 2009a, 2014)	120	25-45	nd	nd	5+	5+	+ to 3+	nd	nd	nd
H2N3 swine/MO/2006 (Chen et al. 2014)	64	24-86	nd	nd	2+	5+	2+ to 5+	nd	nd	nd
H5N1 VN/1203/2004 (Suguitan et al. 2006)	12	14	15	10-12	3+	3+	-	4+	5+	5+
H5N1 HK/213/2003	466	10	1,955	14-24	5+	5+	-	5+	5+	5+
H6N1 teal/HK/W312/97 (Chen et al. 2009b)	269	20-57	nd	nd	3+	5+	- to +	5+	nd	nd
H7N3 BC CN-6/2004 (Joseph et al. 2008)	160	11	264	10	3+	5+	3+	2+	3+	2+
H7N7 NL/219/2003 (Min et al. 2010)	12	10-15	41	54-80	2+	5+	2+ to 3+	3+	5+	5+

^a The H9N2 ck/HK/G9/97 vaccine virus was not evaluated in ferrets

^b Serum neutralizing antibody titer measured in a microneutralization (MN) assay

^c Efficacy was assessed by reduction of replication of wild-type challenge virus compared to mock-vaccinated animals. <10-fold reduction, + 10 to 99-fold reduction, 2+ 100 to 999-fold reduction, 3+ 1,000 to 9,999-fold reduction, 4+ >10,000-fold reduction, 5+ complete protection from replication

^d Heterologous challenge viruses for H2 vaccines: Japan/57 (H2N2); swine/MO/06 (H2N3); for H5 vaccines: HK/491/97, HK/213/2003; Vietnam/1203/2004, Indonesia/5/2005; for H6 vaccines: Dk/HK/77, mallard/Alberta/85; for H7 vaccines: turkey/V A/55/02 (H7N2), turkey/UT/24721-10/95 (H7N3); chicken/BC/04 (H7N3), Netherlands/219/03 (H7N7), tk/Italy/99 (H7N1), turkey/England/63 (H7N3); for H9 vaccines: HK/11073/99

nd Not done

Protective efficacy of pLAIVs was assessed by determining the ability of one or two doses of intranasally administered vaccine to protect mice from lethal challenge with *wt* virus or to prevent replication of *wt* challenge virus in the URT and LRT of mice and ferrets (Tables 4 and 5). Even vaccines that were poorly immunogenic (e.g., VN04 H5N1 and H7N7 vaccines) provided complete protection from lethal challenge following one dose of vaccine (Min et al. 2010; Suguitan et al. 2006). Two doses of the pLAIVs provided complete protection from replication of homologous *wt* challenge viruses in the URT and LRT of mice and LRT of ferrets; challenge virus titers in the URT of ferrets were reduced compared to mock-immunized animals but protection in the URT was not always complete (Tables 4 and 5). A single dose of vaccine conferred partial to complete protection from challenge virus replication; higher serum antibody titers (MN >80 and HAI of 320) were associated with complete protection. The VN04 H5N1 and H7N7 vaccines that failed to induce a robust antibody response conferred only partial protection following a single dose of vaccine, with a reduction in titer of challenge virus in the range of 40 to 1500-fold compared to mock-immunized animals (Min et al. 2010; Suguitan et al. 2006).

Because it is not possible to predict which strain within a subtype will cause a pandemic and influenza viruses continue to evolve in nature, we evaluated the antibody response and protection from challenge with heterologous viruses. In fact, the data confirmed our hypothesis that a pLAIV would elicit a cross-reactive antibody response against other strains from the same subtype. Two doses of pLAIV elicited cross-reactive antibodies and provided near complete protection from replication of heterologous *wt* viruses in the URT and LRT of mice and LRT of ferrets (Tables 4 and 5).

Another important caveat of preclinical evaluation of pLAIVs is that the studies are generally carried out in influenza-naïve unprimed animals while humans with a prior history of influenza infection or vaccination are immunologically primed. This difference was striking in the H1N1 pandemic in 2009: although people younger than 60 years of age were seronegative to the H1N1pdm virus, all but the youngest children responded to a single dose of inactivated H1N1pdm vaccine while unprimed animals required two doses of inactivated vaccine. We investigated the ability of *wt* seasonal H1N1 virus, seasonal H1N1 LAIV and seasonal TIV to prime mice for an antibody and cellular immune response to a single dose of H1N1pdm LAIV (Chen et al. 2011a) and found that prior exposure to live seasonal H1N1 virus, *wt*, or LAIV, primed mice but seasonal TIV did not. While two doses of H1N1pdm LAIV induced a robust neutralizing serum antibody response, mice that were primed with seasonal H1N1 infection or seasonal LAIV followed by one dose of H1N1pdm LAIV were equally well protected from challenge, in the absence of neutralizing serum antibody (Chen et al. 2011a). Cheng et al. compared TIV and LAIV in ferrets and found that LAIV was superior to TIV in inducing influenza-specific immunity in naïve ferrets (Cheng et al. 2013). Although both types of vaccines induced comparable humoral immune responses in previously primed ferrets, only LAIV provided partial protection from heterologous seasonal H1N1 virus challenge (Cheng et al. 2013). The findings in LAIV-immunized naïve ferrets may explain the efficacy of LAIV in young children.

7.2 *Clinical Studies*

The pLAIVs described above have been evaluated in phase I clinical trials in small cohorts of healthy adults younger than 50 years of age (Karron et al. 2009a, b; Talaat et al. 2009, 2011, 2012). In order to minimize the risk of reassortment between a pLAIV bearing novel HA and NA genes and circulating human influenza viruses, the clinical trials are conducted in an inpatient unit during months when seasonal influenza activity is not detected, typically between April and December. Vaccine recipients are admitted to the inpatient unit a day prior to vaccine administration in case they are incubating an intercurrent respiratory virus infection and also to determine whether they can adjust to a 9–10 day stay in the inpatient unit. The studies are conducted as open-label inpatient trials with all participants receiving vaccine. Vaccine is administered intranasally as a nasal spray except for the H9N2 ca vaccine that was administered as nose drops. Participants are examined daily while on the inpatient unit by a health care provider. After discharge from the inpatient unit, vaccinees are asked to return to the outpatient clinic. At each visit, staff obtain vital signs, review interim histories, and obtain blood and nasal wash (NW) samples for antibody testing (Coelingh et al. 2014).

Vaccine safety is assessed through daily examinations and infectivity is assessed by viral culture and by realtime reverse transcription-polymerase chain reaction testing of NW specimens. Immunogenicity is assessed by measuring HAI antibodies, neutralizing antibodies, and IgG or IgA antibodies to recombinant HA in serum or NW. As reviewed by Coelingh et al. (2014), the pLAIVs were restricted in replication and were variably and generally not immunogenic in terms of antibody responses in humans. These findings were unexpected because it was anticipated that adults receiving pLAIVs bearing novel HA and NA proteins would respond to the vaccines like children receiving seasonal LAIV, shedding vaccine virus for several days, and developing serum and/or NW antibody responses. However, the H5N1 pLAIV appears to have established long term immune memory because when H5N1 pLAIV vaccine recipients were re-called and given a dose of unadjuvanted inactivated subunit vaccine, a rapid robust and high quality antibody response was detected (Talaat et al. 2014). This observation is being explored further in a series of ongoing clinical trials that will provide insights into the durability of pLAIV-mediated immune memory by determining the optimal interval between the LAIV and inactivated vaccine.

8 Mediators and Correlates of Protection

While serum HAI antibody is a correlate of protection for inactivated influenza virus vaccines, it is not an absolute correlate of protection for LAIV. In addition to serum antibody responses, seasonal LAIVs induce mucosal and cell-mediated immunity (Hoft et al. 2011). Although protective efficacy and effectiveness of seasonal LAIV have been repeatedly demonstrated in preclinical and clinical

studies, an immune correlate of protection has not been identified. As reviewed by Bandell et al. (2011), seasonal LAIV was efficacious even in the absence of a rise in serum HAI antibody.

8.1 Studies in Animal Models

The role of humoral and cellular immune mediators and distribution of immune effectors induced by eight different LAIVs at mucosal and systemic sites were evaluated in mice (Lau et al. 2011). All LAIVs tested induced robust systemic immune responses but variable pulmonary immunity. The magnitude of lung immunity, including pulmonary IgA antibody and memory CD8+ T lymphocytes, induced by the vaccines depended on the replication efficiency of the LAIVs and the induction of cytokines/chemokines in the lungs. Both cellular and humoral immunity contribute to the protection provided by LAIV; the relative contribution of the two effector arms in viral clearance depends on the location and the rate of replication of a particular vaccine virus. The relevance of these findings for the human experience was not clear because LAIVs do not replicate in the LRT of humans. Therefore, an upper respiratory tract immunization (URTI) model was developed in mice to mimic the human situation. This permitted assessment of the protective efficacy of an H5N1 LAIV against highly pathogenic H5N1 virus challenge in the absence of significant pulmonary immunity (Lau et al. 2012). The experiments demonstrated that cellular immunity in the lungs is essential for protection against lethal wild-type H5N1 challenge, whereas influenza-specific serum ELISA antibodies and splenic influenza-specific CD8+ CTLs made little contribution to this protection. Optimal protection against wild-type virus challenge requires maturation of humoral responses, with the development of neutralizing activity. Passive transfer of postvaccination serum to naïve mice demonstrated that the magnitude of the humoral response and access of antibodies to the respiratory tract are equally important determinants of protection (Lau et al. 2012).

In a recent study comparing LAIV and TIV in ferrets, while TIV only induced immune responses in primed ferrets, LAIV induced influenza-specific antibody and T cell responses in both naïve and primed ferrets (Cheng et al. 2013). In addition to influenza-specific serum IgA and IgG antibodies, CD4+ and CD8+ T cells were demonstrated in the circulation and in paratracheal lymph nodes and the latter correlated with protection from challenge virus replication in the URT (Cheng et al. 2013).

8.2 Studies in Humans

Immunity to influenza A viruses in humans is conferred primarily by antibodies directed at the HA and NA glycoproteins. Both serum IgG and mucosal IgA antibodies can independently contribute to resistance to influenza virus in humans

(Clements et al. 1986), with serum IgG antibodies providing protection primarily to the LRT and IgA antibodies providing protection primarily to the URT (reviewed in Murphy and Coelingh 2002). Inactivated influenza vaccines induce higher titers of serum antibodies than LAIV in primed individuals, but LAIV are more efficient in inducing mucosal IgA antibody responses (Clements et al. 1986). Thus, serum HAI antibody is not an absolute correlate of protection for LAIV (Ambrose et al. 2008; Edwards et al. 1994; Treanor et al. 1999). Instead, NW IgA induced by LAIV or natural infection is associated with resistance to reinfection. LAIV viruses are shed for a longer duration and to higher titers in immunologically naive, seronegative children than in seronegative adults who have previous experience with homotypic influenza viruses (Murphy and Coelingh 2002). Homotypic immunity induced by prior infection with a drift variant restricts replication of LAIVs in seronegative adults or children to lower titer and for a shorter duration (reviewed in Murphy and Coelingh 2002). In seropositive adults and children, generally only the subset of vaccinees with low preexisting nasal HA-specific IgA antibody titer become infected and they shed a very small quantity of virus.

Although the contribution of CD8+ T cells to protection from influenza virus infection in humans is less clear than in mice and ferrets, the induction of influenza-specific CD4+, CD8+, and $\gamma\delta$ T cells by LAIVs has been demonstrated in children (He et al. 2006; Hoft et al. 2011). In a large field study of seasonal LAIV administered to children, >100 spot forming cells in an interferon-gamma (IFN- γ) ELISpot assay was associated with vaccine efficacy (Forrest et al. 2008). In a prospective study in the UK during the 2009 pandemic, Sridhar et al. found that in the absence of cross-reactive neutralizing antibodies, CD8+ T cells specific to conserved viral epitopes correlated with cross-protection against symptomatic influenza. Higher frequencies of preexisting T cells to conserved CD8 epitopes were found in individuals who developed less severe illness, with total symptom score having the strongest inverse correlation with the frequency of IFN- γ + interleukin-2- CD8+ T cells (Sridhar et al. 2013).

A robust Type I (IFN γ) memory response was observed including production of cytokines (GM-CSF, IL-1 β , IFN α , IL-6) and chemokines (CCL5 and CXCL8 early and CCL2 and CXCL10 late) involved in T cell activation and recruitment, when peripheral blood mononuclear cells from LAIV recipients were cultured with LAIV (Lanthier et al. 2011). Influenza and respiratory syncytial virus infections have been shown to elicit different innate immune signatures by microarray and transcriptomics studies (Herberg et al. 2013), suggesting that LAIVs may also be associated with an innate immune signature. LAIV induced higher expression of type I IFN and interferon stimulated genes (ISGs) than TIV in young children where their genome-wide transcript profiles in whole blood were compared at 7 days following vaccination with LAIV or TIV (Zhu et al. 2010); some of these changes may serve as biomarkers of early responses to LAIV.

9 Future Directions

Live attenuated influenza vaccines are important public health tools for the prevention of seasonal and pandemic influenza. The ability to rapidly produce a vaccine for use in the event of a pandemic requires appropriate infrastructure and capacity that are built on experience with seasonal influenza vaccine. For both seasonal LAIV and pLAIV, the identification of a biomarker that is a reliable immune correlate of protection is a high priority, because this will allow a vaccine against a novel influenza virus to be licensed rapidly in the event of a pandemic threat. The observation that pLAIVs induce long-term immune memory requires further investigation and the feasibility of implementing this immunization strategy needs to be assessed.

Acknowledgments The seasonal influenza vaccine seed development program was conducted by the strain variant team of MedImmune at California. The pandemic vaccine program was conducted under a Cooperative Research and Development Agreement between MedImmune and NIH and was supported in part by the Division of Intramural Research, National Institute of Allergy and Infectious Diseases, NIH. We thank Dr. Zhongying Chen for providing data for Fig. 2 and Drs. Raburn Mallory and Kathy Coelingh for suggestions and comments.

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