Targeting the HA2 subunit of influenza A virus hemagglutinin via CD40L provides universal protection against diverse subtypes

X Fan1,2,9, AM Hashem3,4,5,6,9, Z Chen7, C Li1, T Doyle3,6, Y Zhang2, Y Yi7, A Farnsworth3, K Xu1, Z Li1, R He8, X Li3,6 and J Wang1

The influenza viral hemagglutinin (HA) is comprised of two subunits. Current influenza vaccine predominantly induces neutralizing antibodies (Abs) against the HA1 subunit, which is constantly evolving in unpredictable fashion. The other subunit, HA2, however, is highly conserved but largely shielded by the HA head domain. Thus, enhancing immune response against HA2 could potentially elicit broadly inhibitory Abs. We generated a recombinant adenovirus (rAd) encoding secreted fusion protein, consisting of codon-optimized HA2 subunit of influenza A/California/7/2009(H1N1) virus fused to a trimerized form of murine CD40L, and determined its ability of inducing protective immunity upon intranasal administration. We found that mice immunized with this recombinant viral vaccine were completely protected against lethal challenge with divergent influenza A virus subtypes including H1N1, H3N2, and H9N2. Codon-optimization of HA2 as well as the use of CD40L as a targeting ligand/molecular adjuvant were indispensable to enhance HA2-specific mucosal IgA and serum IgG levels. Moreover, induction of HA2-specific T-cell responses was dependent on CD40L, as rAd secreting HA2 subunit without CD40L failed to induce any significant levels of T-cell cytokines. Finally, sera obtained from immunized mice were capable of inhibiting 13 subtypes of influenza A viruses in vitro. These results provide proof of concept for a prototype HA2-based universal influenza vaccine.

INTRODUCTION

Current seasonal influenza vaccines provide substantial protection by inducing strain-specific neutralizing antibodies (Abs) against the viral hemagglutinin (HA). However, the constantly changing nature of the HA antigenic epitopes presents a daunting challenge to vaccine developers and manufacturers, making it necessary to update the influenza vaccines annually to ensure strains match and thus to achieve sufficient protection in the human population.1–2 Although direct human-to-human transmission of avian influenza viruses has been limited so far, the ability of viruses such as H5N1, H7N7, H7N9, and H9N2 to infect humans and cause disease has raised the concern about their potential to cause devastating pandemics.3–6 All of this underscores the importance of developing a universal influenza vaccine capable of inducing broadly protective immunity against the conserved regions of the viral proteins.

The HA protein represents an attractive target for preventive and therapeutic intervention against influenza because it plays critical roles in the early stages of virus infection by binding to viral receptors on target cells and mediating viral and cellular membranes fusion.7 HA is a type I transmembrane homotrimer protein, in which each monomer is composed of two subunits, HA1 and HA2, linked by a single disulfide bond.7 The HA1 subunit forms a membrane-distal globular head that contains the receptor-binding site and most of the highly variable immunodominant antigenic regions recognized by neutralizing Abs.8,9

1National Institutes for Food and Drug Control, Beijing, China. 2Department of Biopharmaceutics, School of Pharmacy, Fourth Military Medical University, Xi’an, China. 3Centre for Vaccine Evaluation, Biologics and Genetic Therapies Directorate, HPFB, Health Canada, Ottawa, Ontario, Canada. 4Department of Medical Microbiology and Parasitology, Faculty of Medicine, King Abdulaziz University, Jeddah, Saudi Arabia. 5Special Infectious Agents Unit, King Fahd Medical Research Center, King Abdulaziz University, Jeddah, Saudi Arabia. 6Department of Biochemistry, Microbiology and Immunology, University of Ottawa, Ottawa, Ontario, Canada. 7Shanghai Institute of Biological Products, Shanghai, China and 8National Microbiology Laboratory, Public Health Agency of Canada, Winnipeg, Manitoba, Canada. Correspondence: X Li or J Wang (sean.li@hc-sc.gc.ca or wangjz@nicpbp.org.cn)
9The first two authors contributed equally to the work.

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On the other hand, HA2 forms most of the highly conserved stem-like structure, which anchors the globular domain to the viral membrane and contains the viral fusion peptide.\textsuperscript{10,11} Several cross-subtype broadly neutralizing Abs directed against the stem-like domain of the viral HA have been isolated from human antibody libraries, and the structural basis of their binding and neutralization capabilities has been elucidated.\textsuperscript{12–17} However, induction of such Abs by vaccination has been associated with limited success. For example, while different novel constructs based on the structural data obtained from these broadly neutralizing Abs studies were investigated; they mainly conferred subtype-specific immunity with limited heterosubtypic protection.\textsuperscript{18–20} Thus, it would be of great interest to consider molecular adjuvants such as cytokines and costimulatory molecules to enhance immunogenicity and potency of the relatively weak HA2 antigen.

One of the key molecules that regulate humoral and cellular immune responses is CD40 and its ligand (CD40L).\textsuperscript{21–23} CD40 is constitutively expressed on all antigen presenting cells, and CD40L is a type II transmembrane multimer protein mainly expressed by activated CD4\textsuperscript{+} T cells transiently as cell-surface or secreted protein.\textsuperscript{21–23} The critical roles of CD40–CD40L interaction in activation of antigen presenting cells, T cells, and B cells as well as immune response induction have been demonstrated in many studies.\textsuperscript{24–28}

The potential of CD40L as a molecular adjuvant has been also investigated by several groups using different strategies and proven to enhance the immune responses and protection against various pathogens and tumors.\textsuperscript{29–39} Although few reports have employed CD40L to enhance immune response against influenza virus using the full-length HA protein or the highly variable domain of the protein in avian hosts,\textsuperscript{40,41} these studies have mainly focused on immune response induction and only showed partial subtype-specific protection without any evidence of universal protection. Furthermore, although the HA2 subunit is highly conserved among the different subtypes of influenza A viruses and may serve as a “universal” influenza vaccine candidate, it appears to be of low immunogenicity and protective efficacy as Abs induced following immunization with HA protein are found to be mainly targeting the HA1 subunit of the proteins.\textsuperscript{12–20} In this study, we investigated the potential and breadth of CD40L as a targeting ligand and molecular adjuvant in enhancing HA2-induced protective immune response. Here, we report that intranasal delivery of the vectored vaccine expressing the fusion protein of HA2 and CD40L completely protect mice against lethal challenges with multiple mouse-adapted viruses and sera obtained from the immunized mice could substantially inhibit 13 subtypes of influenza A virus.

**RESULTS**

**Construction of rAd vaccines and protein expression**

We generated recombinant adenovirus (rAd) vector expressing the wild-type (wt) or codon-optimized (Opt) HA2 from influenza A/California/7/2009(H1N1) virus fused to a secretion signal (S) at the N terminus and to a trimerized form of murine CD40L ectodomain at the C terminus (rAd-SHA2(wt)FCD40L and rAd-SHA2(Opt)FCD40L, respectively); thus influenza HA2 can be secreted from infected cells and targeted to CD40 receptors on antigen presenting cells (Figure 1a). In addition, we generated rAd vectors expressing non-targeted but secreted monomeric wt HA2 (rAd-SHA2(wt)) or Opt HA2 (rAd-SHA2(Opt)). A rAd encoding a non-targeted but secreted trimeric Opt HA2 was also generated (rAd-SHA2(Opt)F) as well as an “empty” vector control (rAd-Control). Prior to animal experiments, the expression of all fusion proteins SHA2(Opt)FCD40L, SHA2(wt)FCD40L, SHA2(Opt)F, SHA2(Opt), and SHA2(wt) was confirmed in vitro (Figure 1b). Western blot analysis also confirmed that only gene products containing the trimerization motif from rAd-SHA2(Opt)FCD40L, rAd-SHA2(wt)FCD40L,
and rAd-SHA2(Opt)F show bands at low and high molecular weights corresponding to the expected molecular sizes of the monomeric and trimeric proteins. Specifically, the proteins without the trimerization motif encoded by rAd-SHA2(wt) or rAd-SHA2(Opt) were found to be at a size around 26–28 KDa corresponding to monomeric protein only. On the other hand, the protein encoded by rAd-SHA(Opt)F was found to form a low molecular weight protein with a band size of 28–30 KDa as well as a high order molecular weight protein with a size around 85–90 KDa corresponding to monomeric and trimeric proteins, respectively. Similarly, western blot analysis of the proteins expressed by rAd-SHA2(wt)FCD40L and rAd-SHA2(Opt)FCD40L revealed a low molecular weight band corresponding to the monomeric form of the protein and a high molecular weight trimeric protein with size around 150 KDa. In addition, only proteins containing CD40L were detected by anti-CD40L Abs but all proteins were detected by anti-HA2 Abs (data not shown).

**CD40L and codon-optimization enhance the immunogenicity of HA2 protein**

In order to evaluate the effects of using CD40L as an adjuvant and targeting molecule on the induction of HA2-specific immune response, mice were intranasally immunized in a prime-boost regimen with different doses (10⁹, 10⁸, or 10⁷ pfu) of the generated constructs and the immune response was analyzed 2 weeks post primary and secondary vaccination. Although mice immunized with any of the three constructs expressing HA2 generated significant levels of local and systemic anti-HA2 Abs compared with control mice immunized with rAd-Control or phosphate-buffered saline (PBS) in a dose–response manner, the codon-optimized construct resulted in the highest titers of circulatory Abs and mucosal IgA (Figure 2 and Supplementary Figures S1 and S2 online). Specifically, targeting the secreted HA2(wt) via CD40L elicited ≥2fold increase in the levels of mucosal (IgA) and systemic (IgG1 and IgG2a) antibody titer compared with the untargeted rAd-SHA2(wt) control after boosting (Figure 2). In mice immunized with rAd-SHA2(Opt)FCD40L, highly significant levels of nasal IgA and circulatory IgG1 and IgG2a Abs with titers greater than fourfold compared with rAd-SHA2(wt) were also generated. It is also of note that the level of antigen-specific Abs elicited by rAd-SHA2(Opt)FCD40L were significantly more than those induced by rAd-SHA2(wt)FCD40L by ≥2folds (Figure 2). Similarly, highly significant levels of mucosal IgA and circulatory IgG1 and IgG2a were also observed 2 weeks post secondary immunization with a dose of 10⁶ pfu but not 10⁷ pfu of these constructs (Supplementary Figure S1). Furthermore, the levels of anti-HA2 IgG1 and IgG2a Abs were still highly elevated up to 2 months post secondary immunization (Supplementary Figure S2). These results indicate that both CD40L-targeting and HA2 codon-optimization significantly enhanced nasal and circulating Ab responses.

Antigen-specific T-cell responses were also evaluated by ELISPOT to determine the effects of CD40L on cellular immune response 2 weeks post primary and secondary immunization. To our surprise, immunization of mice with rAd-SHA2(wt) did not induce any significant levels of IFN-γ, TNF-α, or IL-4 compared with control groups (rAd-Control or PBS) after priming or boosting (Figure 3 and Supplementary Figure S3). On the other hand, targeting secreted HA2(wt) or HA2(Opt) via CD40L elicited highly significant levels of these cytokines at both time points in a dose–response fashion (Figure 3 and Supplementary Figure S3). Interestingly, there was no significant difference between these two groups suggesting that CD40L preferentially enhanced the induction of cellular

![Figure 2](a) CD40L and HA2 codon-optimization enhance circulating and nasal anti-HA2 antibody response. HA2-specific antibody titers at 2 weeks after (a) priming and (b) boosting are shown for circulating IgG1 and IgG2a, and nasal IgA. Balb/c mice were intranasally immunized with 10⁶ pfu of each rAd virus or with PBS on day 0 and boosted 2 weeks post primary immunization. Data are shown as mean titer ± s.d. from one experiment out of three independent experiments, with n = 3 mice per treatment group in each experiment. **P < 0.0001, ***P < 0.001, ****P < 0.01 and *P < 0.05 (one-way ANOVA with Bonferroni post-test).
Substantially enhance humoral and cellular immune responses. Indicate that the use of CD40L as targeting molecule in a ... were sacrificed at 2 weeks (b) secondary immunization and splenocytes were isolated for IFN-γ, TNF-α, and IL-4 measurement by ELISpot. Data shown as mean numbers of spot-forming cells (SFCs) ± s.d. from one experiment with n = 3 per treatment group. Each sample was tested in triplicates. ****P < 0.0001, ***P < 0.001, **P < 0.01 and *P < 0.05 (one-way ANOVA with Bonferroni post-test).

Figure 3 CD40L enhances T-cell responses to HA2. Balb/c mice were intranasally immunized twice with 10^9 pfu of each rAd virus or with PBS. Mice were challenged 2 weeks following boosting by intranasal influenza viruses from different HA subtypes. Immunized mice were sacrificed at 2 weeks (a) post primary or (b) secondary immunization and splenocytes were isolated for IFN-γ, TNF-α, and IL-4 measurement by ELISpot. Data shown as mean numbers of spot-forming cells (SFCs) ± s.d. from one experiment with n = 3 per treatment group. Each sample was tested in triplicates. ****P < 0.0001, ***P < 0.001, **P < 0.01 and *P < 0.05 (one-way ANOVA with Bonferroni post-test).

Immune responses in vaccinated mice. Here, targeting HA2 protein via CD40L always induced higher levels of IFN-γ followed by TNF-α and finally IL-4. Collectively, these results indicate that the use of CD40L as targeting molecule in conjunction with codon-optimization of the antigen could substantially enhance humoral and cellular immune responses.

Targeting secreted HA2 via CD40L confers heterosubtypic protection in vivo

Having observed the enhanced immune responses in mice immunized with CD40L-targeted HA2, we investigated the potential of these vaccines to provide cross-protection against influenza viruses from different HA subtypes. Immunized mice were challenged 2 weeks following boosting by intranasal administration with 5 × LD50 of mouse-adapted influenza A/Puerto Rico/8/1934(H1N1), A/Jingke/30/95(H3N2) or the avian A/chicken/Jiangsu/7/2002(H9N2) strains. As shown in Figure 4, although immunization of mice with rAd-SHA2(wt) at a dose of 10^9 pfu provided 50–60% protection against the same HA group 1 viruses, i.e., A/Puerto Rico/8/1934(H1N1), and A/chicken/Jiangsu/7/2002(H9N2) strains, only 30% protection was observed against the H3N2 virus, which belongs to HA group 2 (Figure 4). Importantly, the same dose (10^9 pfu) of either rAd-SHA2(wt)FCD40L or rAd-SHA2(Opt)FCD40L conferred complete protection against all the three viruses regardless of the subtype (Figure 4).

Such protective effects were observed in a dose–response fashion as lower amounts of antigens correlated with decreasing levels of protection, with the CD40-targeted vaccines consistently affording better protection than the untargeted HA. Specifically, one log less of the two targeted vaccines still provided complete protection against both H1N1 and the avian H9N2 viruses (Supplementary Figure S4). However, rAd-SHA2(Opt)FCD40L afforded the best protection against the more phylogenetically distant H3N2 virus (60% protection) while rAd-SHA2(wt)FCD40L protected 30% of the mice only (Supplementary Figure S4). The strong protection by rAd-SHA2(Opt)FCD40L was further confirmed by the superior protection against all three viruses compared with rAd-SHA2(wt)FCD40L and rAd-SHA2(wt) when a very low dose of 10^7 pfu was employed for immunization (Supplementary Figures S4 and S5).

We next determined the level of virus titers in the lung and nasal tissues. As shown in Figure 5, immunization with rAd-SHA2(wt)FCD40L and rAd-SHA2(Opt)FCD40L showed ≥ 3 log, 2 log, and 1.5 log reduction in H1N1, H9N2, and H3N2 lung and nasal viral loads compared with control groups, respectively. On the other hand, viral titers in mice immunized with rAd-SHA2(wt) were reduced by ≤ 1.5 log as compared with control groups. Taken together, these data demonstrate that not only CD40L but also HA2 codon-optimization substantially enhanced the protective efficacy of HA2 and significantly promoted heterosubtypic immunity against H1N1 strain as well as the divergent viruses, which belong to the phylogenetically distinct H3 and avian H9 subtypes.

Targeting codon-optimized HA2 via CD40L elicits broadly inhibitory Abs against 13 subtypes of influenza A virus

As all of the vaccine constructs were made using HA2 subunit of H1N1, we next wished to determine whether sera obtained from mice immunized with the various vaccine constructs could inhibit 13 different subtypes of viruses available in our laboratories (H1–13). It needs to be mentioned that the other three subtypes (H14–16) are very rare and also not available to us.4 Here, we used an in vitro microneutralization assay to determine whether the sera from immunized mice could inhibit all the subtypes and whether there is any difference between the
sera obtained from mice immunized with the targeted or untargeted vaccines. As shown in Figure 6, only sera collected from mice immunized with rAd-SHA2(Opt)FCD40L showed potent cross-inhibitory activity against 13 HA subtypes, whereas sera collected from rAd-SHA2(wt)FCD40L-vaccinated mice demonstrated protective activity against most subtypes except for H4 and H5 strains. Surprisingly, antisera collected from rAd-SHA2(wt) immunized mice failed to inhibit any of the tested subtypes. These results clearly underscore the role of CD40L and codon-optimization in enhancing the HA2-based immune response and the induction of broadly inhibitory Abs against all influenza A viruses.
CD40L but not the trimerization motif enhances the breadth of HA2-based protection

Having observed the robust protective effects of rAd-SHA2(Opt)FCD40L against diverse subtypes, we next wished to determine whether the trimerization motif rather than the CD40L is responsible for the enhanced immune response we observed. To this end, rAd-SHA2(Opt) and rAd-SHA2(Opt)F were compared with rAd-SHA2(Opt)FCD40L. As shown in Figure 7a–c, no significant differences in animal survival rate or weight loss were observed between rAd-SHA2(Opt) and rAd-SHA2(Opt)F, suggesting that the trimerization motif itself cannot enhance the protection regardless of the trimeric fusion protein encoded by rAd-SHA2(Opt)F. In contrast, although it did not prevent infection and weight loss, the addition of the CD40L using rAd-SHA2(Opt)FCD40L resulted in 100% protection of the animals against the three influenza subtypes (Figure 7a–c). Moreover, these protection effects correlated well with the antibody and cytokine levels in these immunized mice (Figure 7d–e). Similar results were also observed when mice were immunized with lower doses of each rAd (Supplementary Figure S6). Taken together, these data provide strong supporting evidence on the effects of CD40L as an adjuvant in inducing enhanced immunological protection against diverse strains of influenza A virus.

DISCUSSION

Current influenza vaccines provide substantial protection in humans, however they mainly induce strain-specific
neutralizing Abs. Most of these Abs target the highly variable regions of the globular head in the HA1 subunit; therefore, they cannot induce heterosubtypic immunity. The globular head in the HA1 subunit of the virus is highly immunogenic and therefore elicits high levels of neutralizing Abs, which block viral attachment to host cell receptors. On the other hand, the HA2 subunit that represents most of the HA stem region show high sequence conservation among the different HA subtypes. Therefore, the HA2 region would be a very attractive target to induce Abs of broad neutralizing activity. However, previous studies suggest that the immunogeneity of the HA2 is very weak, and the low levels of the Abs in patients may not be sufficient to provide protection.

In this communication, we described a novel “universal” influenza vaccine based on the HA2 subunit, in which we utilized CD40L, a key linker between the innate and the adaptive immune responses, not only as an adjuvant but also as
a targeting molecule to efficiently stimulate antigen presenting cells and to enhance HA2-specific immune responses. The data presented here clearly demonstrate that using CD40L augmented HA2-specific immune responses and enhanced its protective efficacy. Specifically, although codon-optimization of HA2 seems to induce significantly higher levels of local and systemic anti-HA2 Abs, targeting secreted wild-type or codon-optimized HA2 via CD40L significantly promoted Abs secretion and HA2-specific cytokines production by T cells compared with untargeted HA2. These results also indicate that the secreted fusion protein HA2-FCD40L apparently retain the capability to elicit cell-mediated immune responses as demonstrated by the enhanced HA2-specific cytokine production by T cells in mice immunized with constructs containing CD40L only.

It is noteworthy that although the HA2 gene of influenza A/California/7/2009(H1N1) virus was used to generate our constructs, effective protection was still achieved in animals challenged with H3 virus, which shares ~50% of sequence homology with A/California/7/2009(H1N1) virus. Moreover, the better protection against A/Puerto Rico/8/1934(H1N1) and avian A/chicken/Jiangsu/7/2002(H9N2), particularly at lower doses of the constructs, is in agreement with the fact that the HA2 subunits of these two viruses share higher homology with that of A/California/7/2009(H1N1), i.e., 92% and 65% identity, respectively.

The universality of this prototype vaccine is also supported by results generated from our in vitro experiments in which 13 different subtypes of influenza A virus (H1–13) were tested. Most of the 13 viruses are not mouse-adapted, therefore cannot be used in animal challenge studies. However, antisera from the CD40-targeted immunogens, particularly the rAd-SHA2(Opt)/FCD40L construct but not rAd-SHA2(wt), were able to significantly inhibit all 13 different subtypes. These data collectively suggest that the head domain of HA (mainly HA1 subunit) masks the recognition of the highly conserved regions in the stem of the HA protein (HA2) by the immune system in addition to the low immunogenicity of HA2.43–45 but CD40L as targeting molecule and molecular adjuvant could overcome the low immunogenicity of the HA2 subunit and can lead to universal protection against divergent influenza strains belonging to group 1 and 2 HA subtypes. Of note, antisera collected from rAd-SHA2-immunized mice failed to show any inhibitory effect in vitro regardless to the observed partial in vivo protection and reduced lung virus titers. This could be due to the mononeric form or the improper folding of SHA2 protein expressed by this construct in the absence of the T4 fibritin especially that some BnAb such as the F16 Ab mediates their inhibitory effect via binding to conformational epitopes in the HA stem trimmer.16 However, the data shown in Figure 7 and Supplementary Figure S6 clearly show that immunization with rAd-SHA2(Opt)/F cannot elicit any significant enhancement in protection or immune response induction compared with rAd-SHA2(Opt) although it can be expressed as a trimer. This clearly suggests that using CD40L is the main factor that is enhancing the protection, immunogenicity, and induction of broadly inhibitory Abs.

Furthermore, the in vitro data shown in Figure 6 suggest the induction of broadly inhibitory Abs by CD40-targeted HA2. However, the animal studies clearly demonstrate that immunization of mice by HA2 targeted via CD40L elicited humoral and cellular immune responses and prevented death but not infection (immunized mice lost nearly 20% of their weight upon challenge). This is consistent with the fact that most anti-HA2 stem region Abs do not prevent viral entry but interfere with downstream steps in viral replication1–7 in contrast to strain-specific neutralizing Abs that target the head domain of HA and completely prevent infection at the viral attachment step. In addition, the in vivo protection observed in this study is similar to the non-sterilizing immunity conferred by internal viral proteins such as the nucleoprotein in which cellular immunity as well as non-neutralizing anti-nucleoprotein Abs have been implicated in protection.42,46 Experiments involving passive transfer of antisera or adoptive transfer of T cells from vaccinated mice into naive animals could provide more mechanistic details and elucidate the role and the importance of both arms of the immune system in protection. However, the induction of enhanced antigen-specific T-cell cytokines and Ab responses, the presence of broadly inhibiting Abs and the protection of mice from death upon vaccination with rAd expressing CD40-targeted HA2 compared with other groups do not only indicate that both cellular and humoral immune response are involved but also suggest that protection was dependent on targeting HA2 protein to CD40-expressing cells via CD40L.

In short, the HA2 has the potential of being a candidate “universal” vaccine but is unlikely to induce sufficient protective immune responses. Expression of the HA2 as a secreted form of fusion protein with CD40L could substantially increase the immunogenicity of HA2 and elicit broadly inhibitory Abs against most influenza A subtypes.

METHODS

Cell lines and viruses. Madin–Darby canine kidney cells, HeLa cells, and HEK-293A cells were cultured in complete Dulbecco’s modified Eagle’s medium with 10% heat-inactivated fetal bovine serum as described previously.47 The mouse-adapted influenza A/Puerto Rico/8/1934(H1N1), A/jingke/30/95(H3N2), and A/chicken/Jiangsu/7/2002(H9N2) viruses were obtained from Shanghai Institute of Biological Products. All viruses were propagated at 37°C in the allantoic cavities of 10-day-old embryonated hen eggs for 3 days. Allantoic fluid was clarified by centrifugation, aliquoted, and stored at –80°C. Viruses were titrated in Madin–Darby canine kidney cells and used for viral challenge. Median lethal dose (LD50) for each virus in Balb/c mice were determined using Reed–Muench formula.48

Generation of rAds. Constructs were designed to express a secreted form of wild-type, rAd-SHA2(wt), or codon-optimized, rAd-SHA2(Opt), influenza HA2 subunit (amino acids 345–566) from influenza A/California/7/2009(H1N1) virus (GenBank Acc. no. ACP41105) fused to 23 amino acids (MILLAVLYCLLWSFQTSAGHFPR) from the human tyrosinase signal peptide (GenBank Acc. no. AHH03020) at the N terminus as previously described.49 Other constructs, rAd-SHA2(wt)/FCD40L and rAd-SHA2(Opt)/FCD40L, were also generated to express SHA2(wt) and SHA2(Opt) fused to a 27 amino-acid fragment from the bacteriophage T4 fibritin trimerization motif (GYIPEAPRDQAYVXRKDEWVLSTFL) connected to the
Recombinant viruses were purified twice by cesium chloride density-gradient centrifugation. The resulting recombinant plasmids were XhoI-linearized and transfected into HEK-293A cells to package the rAd vectors. Recombinant viruses were purified twice by cesium chloride density-gradient centrifugation, titrated, and stored at -80°C until used. Cloning was confirmed by Xhol digestion and PCR. Viruses were titrated by plaque assay in HEK-293A cells.

**Protein expression in cell culture.** The generated rAds were used to infect confluent HeLa cells in six-well plates at a multiplicity of infection of 100. Forty-eight hours later, protein expression in cell lysates of infected cells was confirmed by western blot of proteins separated on sodium dodecyl sulfate-polyacrylamide gel electrophoresis or native sodium dodecyl sulfate-polyacrylamide gel electrophoresis as previously described using anti-CD40L monoclonal Abs and rabbit anti-HA Abs (Sino Biological Inc., Beijing, China).

**Animal study.** Six to eight-week-old female Balb/c mice were maintained in a pathogen-free animal facility at Shanghai Institute of Biological Products. All animal experiments were conducted in accordance with Institutional Guidelines and with the approval of the Animal Care and Use Committee. Mice were intranasally immunized twice with 10^6 pfu, 10^7 pfu, or 10^8 pfu of each rAd construct (rAd-SHA2(Opt)ECD40L, rAd-SHA2(wt)ECD40L, rAd-SHA2(Opt)EpAdxsi, rAd-SHA2(Opt), rAd-SHA2(wt), or rAd-Control) and used for microneutralization. Approximately 100 TCID50 of each virus (viruses used were previously described) were mixed with an equal volume of serially diluted sera pooled from three mice in each group in 96-well plates and incubated for 1 h at 37°C. Virus-Ab mixture was transferred onto confluent monolayers of Madin–Darby canine kidney cells and incubated at 37°C for 20 h. Cell monolayers were washed twice with PBS, fixed with ice-cold 80% acetone for 10 minutes and viral nucleoprotein was detected by indirect ELISA using rabbit anti-nucleoprotein Abs (Sino Biological).

**Data analysis.** Statistical analysis was conducted using either one-way or two-way analysis of variance when appropriate. Bonferroni post-test was used to adjust for multiple comparisons between the different groups. All statistical analysis was conducted using GraphPad Prism software (San Diego, CA).

**SUPPLEMENTARY MATERIAL** is linked to the online version of the paper at http://www.nature.com/mi

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**DISCLOSURE**

The authors declared no conflict of interest.

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