Vaccine-induced antibodies target sequestered viral antigens to prevent ocular HSV-1 pathogenesis, preserve vision, and preempt productive neuronal infection

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The cornea is essential for vision yet highly sensitive to immune-mediated damage following infection. Generating vaccines that provide sterile immunity against ocular surface pathogens without evoking vision loss is therefore clinically challenging. Here, we tested a prophylactic live-attenuated vaccine against herpes simplex virus type 1 (HSV-1), a widespread human pathogen that can cause corneal blindness. Parenteral vaccination of mice resulted in sterile immunity to subsequent HSV-1 challenge in the cornea and suppressed productive infection of the nervous system. This protection was unmatched by a relevant glycoprotein subunit vaccine. Efficacy of the live-attenuated vaccine involved a T-dependent humoral immune response and complement C3 but not Fcy-receptor 3 or interferon-α/β signaling. Proteomic analysis of viral proteins recognized by antiserum revealed an unexpected repertoire dominated by sequestered antigens rather than surface-exposed envelope glycoproteins. Ocular HSV-1 challenge in naïve and subunit-vaccinated mice triggered vision loss and severe ocular pathologies including corneal opacification, scar formation, neovascularization, and sensation loss. However, corneal pathology was absent in mice receiving the live-attenuated vaccine concomitant with complete preservation of visual acuity. Collectively, this is the first comprehensive report of a prophylactic vaccine candidate that elicits resistance to ocular HSV-1 infection while fully preserving the cornea and visual acuity.

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INTRODUCTION
Vaccine immunology research classically focuses on generating sterile immunity and identifying the mechanisms responsible for protection against infection. However, this approach is inadequate when considering pathogens that affect delicate organs and tissues such as the eye and nervous system. While the eye is well known as an immune-privileged organ, it remains highly susceptible to inflammatory damage. For this reason, many regulatory mechanisms temper ocular inflammation to preserve visual clarity.1–3 Nonetheless, excessive inflammatory responses in the eye often break tolerance, contribute to permanent vision loss, and significantly impact quality of life.4–6 Clinical management of ocular infections is often challenging and requires close attention to controlling both the pathogen and host inflammation to preserve the visual axis.7,8 Accordingly, it is important to consider the potential of vaccine-induced inflammatory responses during the initial stages of vaccine development when targeting pathogens that commonly affect the eye.

Herpes simplex virus type 1 (HSV-1) is a widespread human pathogen that is of particular relevance to this topic. In addition to being a leading cause of infectious corneal blindness, HSV-1 is a clinically important cause of encephalitis and has recently emerged as the leading cause of primary genital herpes in women of childbearing age in the USA.9–11 The success of the pathogen lies in its ability to evade immune responses and establish latency in sensory neurons for the life of the host. Furthermore, the total reservoir of latent virus in the trigeminal ganglia (TG), which supply sensory innervation to orofacial mucosal sites, correlates with reactivation risk and clinical disease burden in animal models.12,13 Chronic viral reactivation in the human eye is associated with a myriad of clinically important corneal pathologies including scarring, neovascularization, and persistent epithelial defects. Current therapies aim to suppress ocular inflammation with steroids and inhibit viral replication with nucleoside analog drugs, but such interventions do not “cure” the disease. Moreover, recurrences frequently persist even when on long-term, prophylactic treatment with these agents.9 Visual morbidity can be so severe that corneal transplantation may be necessary to restore vision, although this remedy often has diminishing returns due to increased graft rejection rates.14 Novel therapies to block HSV-1 pathogenesis are in development.15–17 Considerable effort has also been applied to developing a therapeutic HSV vaccine to alleviate viral reactivation in patients with recurrent outbreaks.18–20 However, we contend that prophylactic vaccination would be a highly effective strategy to prevent HSV-1-associated disease in the eye, skin, and nervous system.

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Herein, we provide a comprehensive immunologic and ophthalmologic evaluation of the protective efficacy of a prophylactic live-attenuated vaccine for HSV-1. Although humans suffer ocular disease largely as a result of HSV-1 reactivation, immunologically naive mice develop robust, clinically relevant corneal disease following primary infection. Therefore, ocular HSV-1 infection in mice serves as a model to study the dynamics and mechanisms of prophylactic protection from the viewpoints of both viral pathogenesis and immune-mediated tissue damage. Using the eye as a clinically relevant site of HSV-1 infection following prophylactic vaccination, we show that a live-attenuated HSV-1 vaccine drives a T-dependent humoral immune response that elicits sterilizing immunity, limits the establishment of viral latency, and fully preserves the visual axis. Thorough characterization of the latter component is missing from nearly all previous efforts to characterize the “efficacy” of vaccines against ocular HSV-1 infection. Moreover, we identify that many dominant HSV-1 antibody targets are not exposed glycoproteins, but rather sequestered antigens only accessible within intracellular compartments. Our previous work shows that humoral immunity is essential for prophylactic protection against ocular HSV-1 infection through a mechanism involving the neonatal Fc receptor (FcRn) and intracellular complement fixation in outbred CD-1 mice.\textsuperscript{21,22} The current investigation uses the genetic and immunologic tools available with the inbred C57BL/6 strain to build upon our previous studies.

METHODS

Mice and immunization scheme

Inbred C57BL/6 WT (stock # 000664), TCRα\textsuperscript{−/−} (stock # 002116), μMT (stock # 002288), FcyRII\textsuperscript{−/−} (stock # 003171), C3\textsuperscript{−/−} (stock # 003641), and Ai14/Rosa26-tdTomato Cre-reporter (stock # 007914) mice were obtained from the Jackson Laboratory (Bar Harbor, ME). Ifnar\textsuperscript{−/−} mice were bred in house. All animals were housed in specific-pathogen free conditions at the Dean McGee Eye Institute. This study was conducted according to protocols approved by the University of Oklahoma Health Sciences Center animal care and use committee. Mice were anesthetized for all invasive procedures by i.p. injection of ketamine (100 mg/kg) plus xylazine (6.6 mg/kg) and were euthanized by cardiac perfusion with 10 ml PBS for tissue collection. Animals were immunized using a prime-boost regimen via ipsilateral footpad (subcutaneous) and hind flank (intramuscular) injections 3 weeks later as previously described\textsuperscript{21}. Briefly, the immunization dosage for the live-attenuated virus was 1 × 10\textsuperscript{6} PFU of HSV-1 0ΔNLS in 10 μl PBS (prime and boost). The subunit vaccine included 2.5 μg of recombinant truncated gD from HSV-2 adjuvanted with 25 μl Imject alum (Thermo Scientific) and 10 μg monophosphoryl lipid A from Salmonella enterica serovar Minnesota (Sigma) in a 35-μl total volume (primer and boost). Animals were 6–10 weeks old upon primary immunization and were acutely challenged with HSV-1 30 days following the secondary boost. Serum neutralization titers were determined prior to challenge and reflect reciprocal serum dilutions upon 50% reduction of cytopathic effect at 24 h using the median tissue culture infectious dose (TCID\textsubscript{50}) for HSV-1 McKrae in vero cell monolayers as previously reported.\textsuperscript{21}

Virus strains and ocular infection

As previously documented,\textsuperscript{21–23} the HSV-1 0ΔNLS vaccine is a live-attenuated recombinant virus derived from the HSV-1 KOS strain; the vaccine strain is rendered attenuated by deletion of the nuclear localization signal on the virally-encoded ICP\textsubscript{0} gene. Unless otherwise indicated, all mice were acutely challenged with 1 × 10\textsuperscript{6} PFU HSV-1 McKrae per eye following partial epithelial debridement of the cornea. Ai14/Rosa26-tdTomato reporter mice were acutely infected with 1 × 10\textsuperscript{5} PFU of transgenic HSV-1 (SCT6 strain) expressing Cre recombinase to visualize productively infected cells in cryosectioned TG by confocal microscopy.\textsuperscript{24} Virus stocks were propagated and maintained in high-titer aliquots as previously described.\textsuperscript{21,24,25}

Assessments of viral pathogenesis

Titers of infectious virus were determined in clarified tissue homogenates by standard plaque assay on monolayers of Vero cells (American Type Culture Collection). Viral lytic gene expression and analysis of latent HSV-1 was evaluated by PCR as previously described.\textsuperscript{26} Briefly, DNA was isolated from TGs at day 30 p.i. and subjected to quantitative PCR for HSV-1 genome using a proprietary primer-probe copy number kit (Primer Design). For relative expression of HSV-1 genes (LAT, TK, ICP27), tissue was harvested at the indicated times p.i. for RNA purification, cDNA conversion, and viral transcript expression was evaluated by real-time PCR. Antiviral gene expression was measured by real-time PCR using PrimePCR technology (Biorad).

Antibody repertoire analysis

To analyze the repertoire of viral proteins recognized by HSV-1 0ΔNLS antiseraum, 4 × 10\textsuperscript{5} Vero cells were plated in each well of a 6 well plate and infected with 4 × 10\textsuperscript{5} PFU HSV-1 McKrae (MOI = 1.0). At 18 h p.i., cells were collected, washed in PBS, and pelleted by centrifugation at 300 × g for 5 min. Pellets were re-suspended in 500 μl of 1% Triton-X100 detergent in PBS (cell lysis buffer) to lyse cells, vortexed every 5 min and placed on ice for a total of 15 min. Cell lysates were clarified by centrifugation at 10,000 × g for 10 min. Next, supernatants from infected and uninfected Vero cells were mixed with 4 μl of serum from naive or immunized mice and 100 μl of immumomagnetic protein G microbeads (Milenyi Biotec) and incubated at 4 °C for 30 min. Microbead/antibody/protein complexes were then immobilized in μMACS magnetic columns (Milenyi Biotec). After thorough column washing with cell lysis buffer, retained proteins were eluted with 500 μl of 100 mM glycine, pH 2.5 and stored at 4°C. Tryptic digests were prepared and analysed by liquid chromatography-tandem mass spectrometry (LC-MS/MS) to identify viral proteins as described in the supplementary methods.

Flow cytometry, adoptive transfers, and in vivo receptor neutralization

Spleens were harvested and single cell suspensions created by filtration through 70 μm mesh in RPMI1640 culture media containing 10% fetal bovine serum and antibiotics. Splenocytes were pelleted and erythrocytes lysed using 0.84% NH\textsubscript{4}Cl (J.T. Baker) in H\textsubscript{2}O. For T cell repertoire profiling, splenocytes were labeled with anti-CD45, anti-CD3, and anti-CD8 antibodies (eBioscience) and MHC class I K\textsubscript{b} tetramers provided by the NIH Tetramer Core Facility for identification of HSV-1 specific CD8\textsuperscript{+} T cells. Tetramer-labeled cells were analyzed using a MacsQuant flow cytometer (Milenyi Biotec) as previously described.\textsuperscript{23} For adoptive transfers into Ifnar\textsuperscript{−/−} mice, bulk preparations of CD3\textsuperscript{+} splenocytes were obtained by immunomagnetic isolation using anti-CD3 microbeads (Milenyi Biotec) according to the manufacturer’s directions. Alternatively, splenocytes were labeled with anti-CD4 or anti-CD8 antibodies (eBioscience) and sorted using an 53e cell sorter (Biorad) for adoptive transfers into TCRα\textsuperscript{−/−} mice. Adoptive transfer of isolated cells was mediated by intravenous retroorbital injection. In some experiments, TCRα\textsuperscript{−/−} mice were injected i.p. with 250 μg anti-mouse CD154 (MR1 clone) or Armenian hamster IgG isotype (both from BioXcell) at days 0, 3, and 6 p.i. to evaluate CD40-dependent immune activation following adoptive transfer of CD4 T cells.

Corneal Pathology

Corneolimbal buttons were harvested from eyes at day 30 p.i., fixed in 4% paraformaldehyde, and permeabilized in 1% Triton-
X100 in PBS to facilitate imaging. Corneas were subsequently labeled with antibodies for Lyve1 and CD31 (Millipore) and flat-mounted for analysis of neovascularization using an Olympus FV1200 laser scanning confocal microscope and 10x objective, as described.21 The total area positive for vasculature per field of view (4 quadrants/cornea) was quantified using Metamorph software (Molecular Devices, Inc.). Alternatively, corneas were assessed for collagen remodeling via multiphoton SHG microscopy using a Leica SP8 confocal microscope equipped with an ultrafast Ti:Sapphire multiphoton Laser (Coherent Chameleon Vision II) as described.26 Briefly, images were generated using an 880 nm wavelength excitation beam on conoelminal buttons suspended in 35 mm glass-bottom dishes (MatTek Corp.) containing 50% glycerol in PBS. Resultant forward and backward SHG signal images from the central cornea were acquired using a 25x water immersion objective lens. Image stacks were qualitatively evaluated for collagen remodeling using a 3-tier scale to reflect normal structure, small regional defects, or widespread scarring. Ranks were empirically determined using images from healthy and scarred corneas for reference. Additionally, a masked clinician monitored corneal opacity using a Kowa SL14 portable slit lamp biomicroscope. Opacification was rated on a scale of 1–6, where 1 indicates stromal haze, 2 indicates moderate opacity, 3 indicates moderate opacity with regional dense opacity, 4 indicates diffuse opacity, and 5 indicates diffuse dense opacity with corneal ulcer as described previously.21 Corneal mechanosensation was measured using a handheld Cochet-Bonnet esthesiometer to assess blink reflexes in alert mice (Luneau Technology). Spatial visual acuity was monitored by optokinetic tracking (OKT) behavioral responses using an OKT apparatus and Optometry software (Cerebral Mechanics, Inc.) as described.27,28 Briefly, masked observers monitored animals centered in a virtual 3D environment for head turning behavior in response to varying frequencies of rotating vertical lines. Visual acuity for left and right eyes is represented by the highest frequency for which animals track clockwise and counter-clockwise line rotation, respectively.

Statistical analysis
Graphpad Prism 5 was used for statistical analysis. Unless indicated otherwise, data shown reflect means ± standard errors of the means. Statistical tests utilized for data analysis are described in each figure legend. Thresholds for significant findings are denoted as: *P < 0.05; **P < 0.01; and ***P < 0.001 in each figure.

RESULTS
The HSV-1 0ΔNLS vaccine requires B and T cells but not IFNα/β signaling for prophylactic protection against HSV-1 neurovirulence. The immunologic compartments required for prophylactic protection against ocular HSV-1 infection were investigated using wildtype (WT) and immune-deficient C57BL/6 mice immunized with a previously characterized live-attenuated virus termed wildtype (WT) and immune-deficient C57BL/6 mice immunized with HSV-1 0ΔNLS survived the challenge (Fig. 1a). Vaccine-mediated protection in Ifnar1−/− mice was unexpected (Fig. 1a) given the central role of IFNα/β in acute antiviral defense against HSV-1.30–32 Immunologically naive mice (WT and Ifnar1−/−) lacked detectable neutralizing antibody to HSV-1 prior to challenge. However, protection in HSV-1 0ΔNLS-vaccinated Ifnar1−/− mice was associated with high pre-challenge serum neutralization titers (Fig. 1b).

The pathogenesis of HSV-1 in naive hosts involves primary infection at mucosal sites with subsequent spread to tissue-innervating sensory nerves where the virus persists indefinitely. To determine the prophylactic impact of each vaccine on acute viral pathogenesis, HSV-1 titers were measured following ocular challenge in the tear film, cornea, and trigeminal ganglia (TG)—the sensory ganglion that supplies innervation to orofacial mucosal sites. Viral shedding was eliminated in the tears of HSV-1 0ΔNLS-vaccinated WT mice by days 3–5 post infection (p.i.), although shedding was sustained through day 5 in all other groups (Fig. 1c). The HSV-1 0ΔNLS vaccine facilitated a 5-log reduction in viral shedding in the tear film of immunized Ifnar1−/− mice relative to naive controls (Fig. 1c). Viral titers were also reduced in corneas from HSV-1 0ΔNLS-immunized WT and Ifnar1−/− mice compared to their respective naive counterparts (Fig. 1d, left). Moreover, infectious virus was not detected in the TG of HSV-1 0ΔNLS-immunized WT or Ifnar1−/− mice at day 5 p.i. (Fig. 1d, right). In contrast, HSV-1 disseminated to the TG of HSV-1 0ΔNLS-immunized μMT mice (Fig. 1e) suggesting that antibody abrogates viral neurovirulence.

Viral neuroinvasion and latency were subsequently investigated using highly sensitive molecular genetic approaches. First, HSV-1 DNA was analyzed by quantitative PCR in the TG of challenged animals at day 30 p.i., a time-point when viral latency is stably established.34 Both WT and Ifnar1−/− mice immunized with HSV-1 0ΔNLS had up to 1000 times fewer HSV-1 genome copies than surviving naive or gD-2-immunized WT animals (Fig. 2a). Virus-encoded latency associated transcripts (LAT) epigenetically regulate neuronal latency.35 Therefore, LAT expression was measured at day 30 p.i. by semiquantitative RT-PCR as a secondary correlate of total latent HSV-1 in the TG. LAT expression was markedly less in TG from HSV-1 0ΔNLS-immunized WT mice compared to gD-2-immunized or naive controls (Fig. 2b). Although the HSV-1 0ΔNLS vaccine limited neuroinvasion and the total amount of latent virus, vaccination did not completely preempt infection of nerve ganglia. Therefore, whether latent virus detected in animals vaccinated with HSV-1 0ΔNLS reflected productive or abortive neuronal infection remained to be determined.35 Accordingly, a dually transgenic model system was utilized to identify nerves with a history of productive HSV-1 infection. For these studies, mice encoding a ubiquitous Cre-inducible tdTomato fluorescent reporter construct were challenged with transgenic HSV-1 encoding Cre recombinase under the infected cell protein ICP0 lytic gene promoter.26 In this system, any surviving cell with a history of viral lytic gene (ICP0) activity will express tdTomato. Trigeminal ganglia from naive and HSV-1 0ΔNLS-vaccinated reporter mice were harvested at day 30 p.i., cryo-sectioned, and imaged by confocal microscopy. Reporter-expressing cells (tdTomato+) were abundant in TG from naive animals but conspicuously absent in TG from HSV-1 0ΔNLS-vaccinated mice following ocular challenge (Fig. 2c). Taken together, our data indicate that the HSV-1 0ΔNLS vaccine drives a T-dependent humoral immune response that rapidly clears HSV-1 from the cornea, impedes neuroinvasion, and blocks productive infection of peripheral nerves following challenge. However, the molecular and cellular mechanisms that mediate protection remained to be determined.
The mechanisms underpinning protection in HSV-1 ΔΔNLS-vaccinated Ifnar1−/− mice were explored further during the acute stage of infection. We have previously shown that FcRn and implicitly intracellular antibody is requisite for ocular protection.22 Because intracellular antibody-pathogen complexes can initiate NFκB, AP-1, and IRF-dependent inflammatory responses,30 we sought to determine whether prophylactic immunity could modulate autophagy-associated or interferon-stimulated gene (ISG) expression in the cornea independent of IFNα/β receptor signaling. Vaccination generated strong serum neutralizing antibody responses (>1:10000) and limited corneal edema in Ifnar1−/− mice following challenge (Figs. 3a, b). Viral replication indicated by thymidine kinase (TK) expression was reduced substantially in the corneas of HSV-1 ΔΔNLS-vaccinated Ifnar1−/− mice relative to naive Ifnar1−/− controls at 48 h p.i. (Fig. 3c). Moreover, TK expression in the corneas of HSV-1 ΔΔNLS-vaccinated Ifnar1−/− mice was similar to levels observed in naive WT controls (Fig. 3c). However, unlike naive WT controls, no viral TK expression was detected in TG from HSV-1 ΔΔNLS-vaccinated Ifnar1−/− mice (Fig. 3d). This data further supports our finding that vaccination with HSV-1 ΔΔNLS prevents productive neuronal infection. Antibody-dependent, Ifnar1-independent signaling did not modulate host antiviral gene expression in the corneas of vaccinated Ifnar1−/− mice aside from upregulation of tetherin (Bst2), inducible nitric oxide synthase (Nos2), and MHC class-I (H2-K1) with concomitant suppression of IL-1β (Figs. 2e, f). Nonetheless, prophylactic protection elicited by HSV-1 ΔΔNLS compensates for a complete loss of the antiviral IFNa/β receptor-signaling pathway.

Cell-mediated immunity conferred by the HSV-1 ΔΔNLS vaccine offers inadequate protection against viral pathogenesis in the absence of antibody

Over the past decade, the HSV vaccine research community has keenly focused on memory T cell responses as a component of prophylactic protection.37-39 While cell-mediated immunity is essential for control of HSV-1 in naive hosts, the extent to which primed T cells contribute to prophylactic protection against HSV-1 remains unclear. Because CD8 T cells maintain HSV-1 latency in the TG,40 we first assessed the CD8 T cell repertoire in immunized mice using MHC class-I tetramers reflecting the top three immunodominant HSV-1 epitopes in C57BL/6 mice.31 By day 5 p.i., the systemic pool of tetramer-specific CD8 T cells surveyed in the spleen was no different among naive, gD-2-immunized, or HSV-1 ΔΔNLS-immunized WT mice (Fig. 4a). Given the modest reduction in HSV-1 titers measured in TG from HSV-1 ΔΔNLS-vaccinated μMT mice compared to those receiving the gD-2 subunit (Fig. 1e), we also surveyed the HSV-specific CD8 T cell repertoire in μMT mice. However, the HSV-specific CD8 T cell repertoire in μMT mice was similar to that measured in WT and, likewise, was not appreciably impacted by vaccination (Fig. S1).

In a previous investigation using outbred CD-1 mice, we identified increased acute T cell infiltration into the corneas of HSV-1 ΔΔNLS-immunized animals relative to naive controls at day 5 p.i.21 Importantly, corneal T cell infiltration is associated with tissue pathology and visual morbidity over time.42-44 Therefore, prophylactic vaccination might enhance effector T cell responses in infected tissues without expanding the repertoire of HSV-specific T cells. Although early enhancement of corneal T cell infiltration was observed in HSV-1 ΔΔNLS-vaccinated animals,21...
humoral immunity elicited by HSV-1 ΔΔNLS requires complement C3 but not FcγRIII for optimal protection against viral neuroinvasion and latency

We have previously shown that acute viral clearance from the tear film of HSV-1 ΔΔNLS vaccinated animals is mediated by complement C3 and not Fcγ receptor 3 (FcγRIII) following ocular HSV-1 challenge.22 To substantiate the effector mechanisms responsible for humoral protection in the cornea and nervous system, HSV-1 pathogenesis was evaluated further in immunized C3−/− and FcγRIII−/− mice. All vaccinated animals exhibited similar pre-challenge serum neutralization titers22 and survived the ocular challenge (Fig. 5a). In terms of latent viral burden at day 30 p.i., HSV-1 ΔΔNLS-vaccinated FcγRIII−/− mice were no more susceptible than vaccinated WT mice (Fig. 5b). In contrast, the amount of latent viral DNA detected in vaccinated C3−/− mice was comparable to naive controls (Fig. 5b).

In order to determine if the dynamics of viral clearance may ultimately impact the magnitude of viral latency, we subsequently assessed viral replication in naive and vaccinated C3−/− mice during acute infection. Viral lytic gene expression was elevated in corneas from HSV-1 ΔΔNLS-vaccinated C3−/− mice relative to vaccinated WT mice by 24 h p.i. (Fig. 5c). By day 3 p.i., lytic gene expression was reduced in corneas and TG from HSV-1 ΔΔNLS-vaccinated WT and C3−/− mice relative to their naive counterparts (Figs. 5d, e). However, there was a trend (p = 0.15) in higher viral lytic gene expression in TG from vaccinated C3−/− animals compared to vaccinated WT (Fig. 5e). Infectious virus was cleared from the corneas of HSV-1 ΔΔNLS-vaccinated WT and C3−/− mice by day 5 p.i. (Fig. 5f). Viral titers were also substantially reduced in the TG of vaccinated C3−/− mice compared to naive controls at day 5 p.i. (Fig. 5g). Nonetheless, low levels of infectious HSV-1 were detected in TG from some HSV-1 ΔΔNLS-vaccinated C3−/− mice at day 5 p.i. (Fig. 5g). Taken together, our data indicate that C3 is categorically essential not only for efficient viral clearance at the point of mucosal exposure22 but also for shielding against the establishment of neuronal latency in animals prophylactically vaccinated with HSV-1 ΔΔNLS.

The HSV-1 ΔΔNLS vaccine elicits antibody responses against heterogeneous viral proteins

Nearly all HSV vaccines tested in clinical trials to date have attempted to elicit humoral responses against exposed virion envelope glycoproteins.11,19,47 We have previously shown that HSV-1 ΔΔNLS antiserum recognizes a broad array of HSV-1 proteins by western blot.21 Here, mass spectrometry was utilized.
to characterize the repertoire of viral proteins recognized by HSV-1 ΔNLS antiserum. Antiserum from HSV-1 ΔNLS-vaccinated WT mice selectively and reproducibly immunoprecipitated twenty HSV-1 proteins from infected cell lysates (Figs. 6a, b). These targets included non-structural infected cell proteins (ICPs), virion surface glycoproteins, and sequestered proteins from the virion tegument and capsid (Fig. 6b). Viral proteins were identified by matching derivative tryptic peptides to the HSV-1 reference proteome as described in the supplementary methods. Envelope glycoproteins only represented 26% of all viral proteins immunoprecipitated with HSV-1 ΔNLS antiserum based on total peptide counts. Non-structural ICPs and tegument proteins each accounted for 10% of recognized viral proteins, and the majority of targets (53%) were identified as capsid proteins (Fig. 6b, inset). Target protein mass likely did not bias results, as no correlations were identified between protein size (kD) and matched tryptic peptide counts (Fig. 6c). Non-viral proteins identified with serum from naive and vaccinated mice likely represent Vero cell xenoantigens or contaminants. Moreover, no viral peptides were identified in serum-free preparations.

Studies of the intraviral "interactome" of HSV-1 have uncovered a complex array of protein-protein interaction networks. Accordingly, it remains possible that the immunoprecipitation products detected by mass spectrometry with HSV-1 ΔNLS

Fig. 3 Contributions of type 1 interferon signaling to prophylactic protection. Type 1 interferon (IFNα/β) receptor-deficient (Ifnar1−/−) mice were prophylactically vaccinated in the footpad with 1 × 10⁵ plaque forming units (PFU) of HSV-1 ΔNLS and ocularly challenged 30 days later with 1 × 10⁵ PFU HSV-1 McKrae per eye. a Spectral domain optical coherence tomography (SD-OCT) imaging of the anterior eye of Ifnar1−/− animals 72 h p.i. b Serum neutralizing titers in vaccinated and naive Ifnar1−/− mice. Relative expression of HSV-1 thymidine kinase (TK) in the cornea (c) and trigeminal ganglia (d) of Ifnar1−/− mice at 48 h post-infection (p.i.). Expression data was normalized to glyceraldehyde 3-phosphate dehydrogenase (GAPDH) expression and relative to tissue from uninfected WT C57BL/6 mice. e Heat map of antiviral gene expression at 48 h p.i. generated using Biorad PrimePCR technology and the National Cancer Institute’s online Cluster Image Map tool (http://discover.nci.nih.gov/cimminer/). Host gene expression was relative to the geometric mean of beta actin, GAPDH, and phosphoglycerokinase 1 expression and normalized to tissue from uninfected WT C57BL/6 mice. Data sets in which significant differences (Student’s T tests) were identified in host gene expression between naive and vaccinated Ifnar1−/− mice are shown in (f). Data in each panel reflect the summary of n = 5 mice/group with two independent experiments. Gene expression levels detected in HSV-infected WT mice are depicted for reference in panels c, d, and f. Abbreviations: TCID₅₀, median tissue culture infectious dose; ND not detected, WT wild-type, Nos2 inducible nitric oxide synthase, Irf1b interleukin 1β, Ifn interferon, Becn1 beclin1, Ripk1 receptor-interacting serine/threonine-protein kinase 1, FcRn neonatal Fc receptor, Ifi interferon regulatory factor, Nfkβ nuclear factor kappa-light-chain-enhancer of activated B cells, Rnasel ribonuclease L, Trim21 tripartite motif-containing protein 21, Eif2ak2 eukaryotic translation initiation factor 2-alpha kinase 2, Bst2 bone-marrow stromal antigen 2 (tetherin), Tlr toll-like receptor, H2-K1 histocompatibility 2, K1 region (MHC class I), Ccl2 C-c motif chemokine ligand 2

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antiserum contained protein-protein complexes that may have contributed to indirect 'false-positive' target identification. None-the-less, twenty viral proteins were consistently identified in our studies despite the hundreds of possible "intraviral" protein-protein interactions confirmed for HSV-1 (Fig. 7d). To explore whether the sequestered viral proteins we identified are probable antibody targets or indirect co-immunoprecipitation products of envelope glycoproteins, we used in silico bioinformatics to query all primary binding partners of gB, gC, gD, and gE with HVint—an open-access HSV-1 interactome database (http://topf-group.ismb.lond.ac.uk/hvint/). Only 7 out of 28 confirmed non-homodimeric envelope glycoprotein-binding partners (25%) were included in our list of sequestered viral proteins identified by proteomic screening. Collectively, our data suggest that effective immunity to HSV-1 provided by prophylactic vaccination with HSV-1 0ΔNLS involves broad antibody responses against both exposed virion

Fig. 4 Contributions of cell-mediated immunity to prophylactic protection. a Flow cytometric analysis of HSV-1 specific CD8 T cells in the spleen of naive and vaccinated WT mice at day 5 p.i. based on tetramer labeling for the top three immunodominant HSV-1 epitopes in C57BL/6 mice including glycoprotein B (gB), and infected cell proteins 6/8 (ICP6/8). Antigen peptide sequences are listed parenthetically (n = 4 vaccinated, 2 naive mice/group; 2 independent experiments). b Viral shedding in the corneas of Ifnar1−/− mice receiving 1 × 10⁶ PFU HSV-1 McKrae per eye (n = 3 mice per group, 2 independent experiments). c Viral titers in the corneas, TG, and brainstem of Ifnar1−/− mice as in (b). d Survival of TCRα−/− mice ocularily challenged with 1 × 10⁶ PFU HSV-1 McKrae/eye 7 days after adoptive transfer of 1 × 10⁶ CD4, 1 × 10⁶ CD8, or 1 × 10⁶ of both CD4 and CD8 T cells i.v. from the spleens of HSV-1 0ΔNLS-vaccinated WT mice concurrent with ocular HSV-1 infection at 1 × 10⁶ PFU HSV-1 McKrae per eye (n = 3 mice per group; 2 independent experiments). e Serum neutralization titers in TCRα−/− mice that survived ocular HSV-1 infection upon receiving T cells from HSV-1 0ΔNLS-vaccinated WT mice (n = 6–15 mice/group; 3 independent experiments). f Serum neutralization titers in TCRα−/− mice that survived ocular HSV-1 infection upon receiving T cells from HSV-1 0ΔNLS-vaccinated WT mice (n = 3–4 mice/group; 2 independent experiments). g Viral titers in the corneas, TG, and brainstem of TCRα−/− mice as in (c), with the addition of naive WT donors (n = 5–6 mice/group; 2 independent experiments). h Survival of TCRα−/− mice ocularily challenged with 1 × 10⁶ PFU HSV-1 McKrae/eye 7 days after adoptive transfer of 1 × 10⁶ CD4 T cells from HSV-1 0ΔNLS-immunized or naive WT mice; TCRα−/− mice were treated with 250 µg anti-CD154 antibody or Rat IgG control i.p. on day 0, 3, and 6 p.i. (n = 8–12 mice + T cells, n = 4 no-cell controls; 3 independent experiments). Data in panels A, C, F were analyzed by one-way ANOVA with Newman–Keuls multiple comparisons tests; data in panels B and E were evaluated two-way ANOVA and Student’s T test, respectively.
envelope glycoproteins and sequestered viral antigens. Moreover, our recent identification that FcRn is essential for humoral protection against ocular HSV-1 infection combined with the identification of multiple sequestered/intracellular viral protein targets herein builds a strong argument that humoral immunity against HSV-1 is not restricted to the extracellular space. While further investigation is ultimately required to determine epitope specificity and binding kinetics, our work underscores that vaccines targeting viral surface antigens alone may be insufficient for protection against HSV-1.

Prophylactic protection elicited by the HSV-1 0ΔNLS vaccine preserves the visual axis against immunopathology and vision loss. The HSV-1 0ΔNLS vaccine amplifies host immunity to mediate protection against HSV-1 pathogenesis, but its impact on visual health remained to be determined in C57BL/6 mice. Therefore, a comprehensive histologic and functional evaluation of corneal health was conducted to establish the protective efficacy of HSV-1 0ΔNLS against ocular immunopathology and vision loss in WT mice. Animals were evaluated for corneal neovascularization, scarring, opacity, and sensory nerve defects at day 30 p.i. Functional visual acuity was tested by optokinetic tracking reflexes. Confocal imaging of corneolimbal buttons revealed that neovascularization was prevented in HSV-1 0ΔNLS-immunized mice (Fig. 7a), whereas robust hemangiogenesis and lymphangiogenesis were observed in corneas from gD-2-immunized and naive mice (Figs. 7b, c). Second harmonic generation (SHG) microscopy showed that central corneal scarring was prevented in nearly all HSV-1 0ΔNLS-immunized mice, although corneas from gD-2-immunized and naive mice exhibited varying degrees of pathologic collagen remodeling (Figs. 7d, e). Additionally, the HSV-1 0ΔNLS vaccine also prevented robust corneal neovascularization and fibrosis in Ifnar1−/− mice (Fig. S2). Consistent with corneal neovascularization and scarring data in WT mice, non-invasive slit lamp biomicroscopy examinations revealed progressive corneal opacification in naive and gD-2-immunized mice following ocular HSV-1 challenge. In contrast, corneal clarity was preserved in HSV-1 0ΔNLS-immunized mice (Fig. 7f). Sensory nerve deficits are another hallmark of corneal herpetic disease, therefore, corneal sensation was measured via Cochet-Bonnet esthesiometry. No sensation loss was detected in HSV-1 0ΔNLS-immunized mice (Fig. 7g). Comparatively, gD-2-immunized and naive mice exhibited moderate to severe sensation loss, (Fig. 7g). Multiple assessments indicate that the HSV-1 0ΔNLS vaccine averts corneal pathology, yet preservation of functional visual acuity remained to be demonstrated. To that end, optokinetic tracking reflexes were monitored as a behavioral indicator of visual acuity. Following challenge, gD-2-immunized
and naive mice sustained significant vision loss (Fig. 7h). However, visual acuity was preserved in HSV-1 0ΔNLS-immunized mice (Fig. 7h). In summary, prophylactic protection elicited by the HSV-1 0ΔNLS vaccine protects not only against HSV-1 pathogenesis but also against resultant inflammatory tissue pathology and visual morbidity.

**DISCUSSION**

Our study demonstrates the prophylactic efficacy of HSV-1 0ΔNLS, a live-attenuated vaccine, against ocular HSV-1 pathogenesis and pathology in C57BL/6 mice. Using multiple approaches, we observed that the HSV-1 0ΔNLS vaccine drives a robust T-dependent humoral immune response against a broad repertoire of HSV-1-encoded proteins to inhibit viral neuroinvasion and latency. While the antiviral IFNα/β signaling pathway and FcγRIII were dispensable for protection, efficacy was diminished in the absence of complement C3. A direct contribution of T cells as antiviral effectors was not observed in our adoptive transfer studies. However, in the case of HSV-1 0ΔNLS, the dominance of antibody-mediated viral clearance may influence the local cytokine milieu to restrain antigen presenting cell activation and ‘inappropriate T cell help’ by favoring regulatory T cell stability and preventing the progression of herpetic keratitis by Th1/Th17-polarized CD4 T cells.50–52 Thorough evaluation of the visual axis is of utmost importance when assessing vaccines for common ocular infections. The concern surrounding this element is underscored by the fact that clinical trials for HSV vaccines have

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**Fig. 6** Identification of viral proteins targeted by vaccine-induced antibodies. a Serum from HSV-1 0ΔNLS-vaccinated and naive WT mice was utilized to immunoprecipitate proteins from HSV-1 infected Vero cell lysates. Precipitated proteins were eluted and analyzed by mass spectrometry. Virus-derived and non-viral proteins were identified by cross-referencing derivative tryptic peptide ions with a reference sequence database. Data reflect the numbers of matched peptides per protein derived from HSV-1 or from other sources immunoprecipitated by HSV-1 0ΔNLS antisera and/or naive serum (n = 5 serum samples/group; 3 independent experiments). Viral proteins are shown in red and non-viral proteins are shown in black. Gray lines indicate a 4-fold change in peptide abundance comparing naive and immune serum. b Repertoire of precipitated HSV-1-derived proteins ranked by average tryptic peptide abundance and labeled according to viral protein class (i.e., non-structural infected cell proteins (ICP), red; capsid, blue; tegument, green or envelope glycoprotein, black). Inset pie chart shows the percentage of viral protein targets by class based on the average number of matched viral peptides (n = 5 serum samples/group; 3 independent experiments). c Pearson correlation analysis was performed on viral targets immunoprecipitated by HSV-1 0ΔNLS antiserum to determine the relationship between the average numbers of matched tryptic peptides and the size of each parent viral protein (kDa). Molecular weights were queried from the HSV-1 reference proteome accessible at [http://www.uniprot.org](http://www.uniprot.org) online. Each data point reflects a viral target protein identified from n ≥ 2 of 5 total serum samples analyzed across three independent experiments. Pearson correlation coefficients (r) and p values were determined using GraphPad Prism. d Graphical representation of the intraviral interactome among the subdivisions of viral proteins. Target proteins identified in the proteomic screen are labeled with a black star. Internal lines show confirmed interactions between HSV-1 proteins. Figure was modified from its original format (Ashford et al. Mol. Cell. Proteomics. 2016 Sep; 15: 2939–2953)48 and reproduced via creative commons 4.0 attribution licensing (http://creativecommons.org/licenses/by/4.0/)
specifically excluded patients with a history of ocular HSV-1 involvement. Furthermore, our comprehensive analysis of the visual axis shows that prophylactic vaccination with HSV-1 ΔNLS effectively controls HSV-1 pathogenesis without eliciting ocular pathology or visual morbidity. These findings are unparalleled by previous studies.

For widespread clinical efficacy, the ideal prophylactic HSV-1 vaccine must preclude the establishment of neuronal latency to mitigate the threat of reactivation-associated shedding, horizontal transmission, and ultimately ocular disease. The exact mechanisms underlying the regulation of HSV-1 latency are not fully understood, yet current knowledge of this process stems from in vitro...
Furthermore, the total amount of latent virus in the TG correlates with reactivation risk in animal models. Recent evidence of lytic gene expression during latency challenges the longstanding stable quiescence model of HSV-1 latency. In immunologically naive animals, HSV-specific CD8+ T cells generated after neuronal infection promote resolution of the lytic cycle and govern neuronal latency. In the absence of prophylactic protection, persistent viral lytic gene activity in neurons during latency contributes to functional exhaustion of tissue-resident antiviral T cells. In contrast, our data show that prophylactic humoral immunity blocks productive infection of neurons and significantly limits the establishment of viral latency following ocular challenge. The cornea is the most densely innervated tissue in the body and a prime target for HSV-1 replication. Accordingly, a vaccine such as HSV-1 0ΔNLS that demonstrably curbs HSV-1 neuroinvasion following corneal infection with a high-titer inoculum as demonstrated herein offers hope in the quest to prevent HSV-1 infection in other sites supplied with fewer sensory nerve fibers.

Although the amount of latent virus in animals vaccinated with HSV-1 0ΔNLS is minimal relative to naive controls, latent virus presents an important safety concern for reactivation potential. Our evidence collectively suggests that the establishment of neuronal latency occurs independent of lytic gene expression (ICP0) in the TG of vaccinated animals. This may reflect delivery of HSV-1 DNA into sensory nerve fibers in peripheral sites without viral replication in neurons (i.e., non-productive infection, the minimal requirement for latency establishment). Accumulating evidence demonstrates the ability of UV-inactivated herpesviruses (HCMV) to deliver viral DNA to cells independent of productive replication—yet the stability of the viral genome is transient in this model. Alternatively, productively infected neurons may be cleared in animals vaccinated with HSV-1 0ΔNLS, but this occurs without obvious sensory deficits. Whether prophylactic protection affects cell-mediated immune responses in neuronal ganglia or HSV-1 reactivation potential remains to be determined. From a clinical perspective, whether a vaccine must completely prevent establishment of herpesvirus latency to be protective against clinical disease is debatable. Longitudinal clinical data show that varicella zoster virus (VZV) reactivation (i.e., “shingles”) is rare in vaccinated children. Moreover, vaccine-induced antibody is the clinical correlate of prophylactic protection against VZV.

Other immunologic aspects requisite for an optimally efficacious prophylactic HSV-1 vaccine remain to be explored. The partial reduction in viral titers in the tear film and TG of HSV-1 0ΔNLS-immunized μMT mice may be attributable to HSV-specific CD4 T cell enhancement of NK cell activity. Furthermore, the necessity of CD4 T cells and CD154 for animal survival may also implicate follicular helper T cells (Tfh) as central players in humoral defense against HSV-1. Whether HSV-1 modulates the function of Tfh cells as an immunoevasion strategy during primary infection in naive hosts is currently unknown. Evidence suggests that strong type 1 IFN responses limit induction of Tfh responses and antibody production during systemic infections. Likewise, whether altered Tfh responses to HSV-1 in naturally infected or prophylactically vaccinated subjects extrapolate into differences in antibody repertoire will need to be determined. Other investigators have focused on prophylactically boosting tissue-resident memory T cell responses to HSV-1 through “prime and pull” approaches in animal models. However, the efficacy of prime and pull against viral dissemination following ocular infection has not been characterized, and the absence of iatrogenic corneal pathology was not convincingly demonstrated. Recruiting T cells to the cornea in an effort to control HSV-1 may not be clinically suitable, as T cells are associated with tissue pathology in HSV keratitis, and progression of inflammatory disease occurs independently of active viral infection.

Effective humoral protection against HSV-1 may involve targeting more than neutralizing exposed surface glycoproteins. Of significant interest is the fact that one of the dominant antigens targeted by HSV-1 0ΔNLS antiserum is HSV-1 gC, a surface glycoprotein that functions to inhibit complement activation. Of equal interest is the observation that a large component of the humoral defense against HSV-1 elicited by prophylactic vaccination with HSV-1 0ΔNLS is targeted against the virion capsid. Future work is necessary to determine if antibody-bound capsids are capable of successfully delivering viral DNA to the nucleus. One caveat to our approach is that the cell lysis buffer utilized may not have dissociated individual capsid protein-protein interactions; nevertheless, a constituent (or constituents) from the capsid were immunoprecipitated by HSV-1 0ΔNLS antiserum. In addition, the UL37 tegument protein functionally associated with neuronal retrograde transport was also recognized by antiserum. Accordingly, it is plausible that intracellular antibody inhibits the function of various viral tegument or transactivator proteins such as ICP4 to reduce infectivity or reactivation potential. It remains to be determined if the ability of antibody to engage sequestered/intracellular antigens is restricted to particular intracellular compartments—and whether intracellular antibody compartmentalization is unique in various cell types. The implications of intracellular humoral protection may shed light on the long-standing efficacy of many other live-attenuated viral vaccines.

Using mass spectrometry to evaluate the repertoire of viral targets recognized by antibodies following immunization with a live-attenuated vaccine encompassing nearly all of the viral proteome may prove to be an effective approach to identify the most relevant antigens for next-generation prophylactic vaccine development. Moreover, identification of sequestered and presumably intracellular proteins that are targeted by antibody supports our recent findings that FcRn arbitrates viral clearance in the cornea by mediating intracellular transcytosis of IgG through the corneal epithelium. Whether effective humoral immunity against HSV-1 can be achieved through targeting a small number of critical antigens or whether a broad repertoire of exposed and sequestered antigens is necessary to achieve clinically meaningful protection in a genetically diverse human population remains to be determined. Nonetheless, the identification of sequestered and intracellular viral proteins as antibody targets reflects a new concept in the pursuit of an effective HSV-1 vaccine. Advancing these concepts forward will require continued investment in improving adjuvants to mediate effective and sustained vaccination.
protection with subunit cocktail vaccines, or increasing acceptance of novel, live-attenuated viruses for clinical utilization.

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AUTHOR CONTRIBUTIONS
D.J.R. planned and conducted experiments, performed data analysis, generated mass spectrometry (Fig. 6); D.M.R. assisted with SHG microscopy (Fig. 7d, Fig. S2); D.J. J.C. performed experiments, edited the manuscript, and supervised all work.

DATA AVAILABILITY
The HSV-1 02NLS vaccine is under US patent protection and distribution would be at the discretion of Rational Vaccines, Inc. Acquisition of the other non-commercial materials would require contact with the original sources listed above.

ADDITIONAL INFORMATION
The online version of this article (https://doi.org/10.1038/s41385-019-0131-y) contains supplementary material, which is available to authorized users.

Competing interests: D.J.C. is an advisory board member of Rational Vaccines, Inc., which has licensed U.S. patents 7758605 and 8802109 involving ICPO-mutant herpesviruses. The remaining authors declare no competing interests.

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