

Activation of NF- κ B in CD8⁺ Dendritic Cells *Ex Vivo* by the γ_1 34.5 Null Mutant Correlates with Immunity against Herpes Simplex Virus 1

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The γ_1 34.5 protein of herpes simplex viruses (HSV) is essential for virulence. Accordingly, an HSV mutant lacking γ_1 34.5 is attenuated *in vivo*. Despite its vaccine potential, the mechanism by which the γ_1 34.5 null mutant triggers protective immunity is unknown. In this report we show that vaccination with the γ_1 34.5 null mutant protects against lethal challenge from wild-type virus via I κ B kinase in dendritic cells (DCs), which sense virus-associated molecular patterns. Unlike mock-treated DCs, DCs primed with the γ_1 34.5 null mutant *ex vivo* mediate resistance to wild-type HSV after adoptive transfer into naïve mice. Furthermore, the γ_1 34.5 null mutant activates I κ B kinase, which facilitates p65/RelA phosphorylation and nuclear translocation, resulting in DC maturation. While unable to produce infectious virus in DCs, this mutant virus expresses early and late genes. In its abortive infection, the γ_1 34.5 null mutant induces protective immunity more effectively in CD8⁺ DCs than in CD8⁻ DCs. This is mirrored by a higher level of interleukin-6 (IL-6) and IL-12 secretion by CD8⁺ DCs than CD8⁻ DCs. Remarkably, inhibition of p65/RelA phosphorylation or nuclear translocation in CD8⁺ DCs disrupts protective immunity. These results suggest that engagement of the γ_1 34.5 null mutant with CD8⁺ DCs elicits innate immunity to activate NF- κ B, which translates into protective immunity.

erpes simplex viruses (HSVs) are human pathogens responsible for a spectrum of diseases, including ocular lesions, genital herpes, and encephalitis (11). HSV infection is also a risk factor in HIV acquisition and transmission. Following infection, HSV initiates a lytic cycle in the mucosal tissues, where viral gene transcription, replication, assembly, and egress ensue. This is followed by a latency in sensory ganglia, where reactivation occurs intermittently, resulting in recurrent infections (11). In this process, HSV interacts with host cells, including dendritic cells (DCs), which capture, process, and present viral antigens. Upon maturation, DCs express high levels of costimulatory molecules. Additionally, DCs release inflammatory cytokines to induce T cell responses that restrict viral infection (1, 54, 68).

DCs detect HSV through multiple pathways (22, 48, 50). For example, plasmacytoid DCs detect HSV through Toll-like receptor 9 (TLR9), whereas conventional DCs sense virus via TLRdependent and -independent pathways (22, 32, 48). It is thought that HSV proteins or RNA intermediates trigger conventional DCs (22, 48). Notably, a complex consisting of HSV glycoproteins B, D, H, and L stimulates DC maturation (50). On the other hand, HSV blocks DC activities (3, 26, 28, 52). In immature DCs, HSV type 1 (HSV-1) downregulates cell surface molecules and cytokines, impairing T cell activation (10, 26, 38, 44, 52). HSV-1 also induces apoptosis of DCs, whereas HSV-2 exerts this activity more rapidly (5, 26, 44). Further, HSV-1 perturbs mature DCs (28, 45), where it induces degradation of CD83 (28, 29). Moreover, HSV-1 reduces levels of chemokine receptors CCR7 and CXCR4, which impedes DC migration (45). Therefore, the interaction of HSV and DCs is a key step that dictates the outcome of viral infection.

We recently reported that unlike wild-type virus, an HSV mutant lacking γ_1 34.5 stimulates DC activation (23). Relevant to this is the fact that the γ_1 34.5 gene encodes a multifunctional protein. In infected cells, HSV-1 γ_1 34.5 precludes the shutoff of protein synthesis mediated by the double-stranded RNA-dependent protein kinase (PKR) (9, 20, 21). It facilitates glycoprotein processing and viral egress (6, 25, 34). The γ_1 34.5 protein also interacts with beclin 1 and inhibits autophagy (40). Additionally, the γ_1 34.5 protein inhibits TANK binding kinase 1 and I κ B kinase in the TLR pathways (24, 61). The precise role of γ_1 34.5 is not fully understood, but its mutation attenuates HSV *in vivo* (4, 8, 33, 35, 43, 51, 56, 63). Of note, vaccination with the γ_1 34.5 null mutant or its derivatives protects from lethal challenge with wild-type virus (41, 46, 55). These studies highlight the γ_1 34.5 null mutant as a potential live attenuated vaccine. Similarly, this idea has been pursued with other HSV mutants. Examples are HSV recombinants lacking gH, gE, ICP0, ICP8, ICP10, or virion host shut-off (vhs) protein (15, 18, 19, 39, 62). While HSV mutants induce antibody and T cell responses upon vaccination (15, 18, 39, 62), the nature of innate immunity remains obscure.

This study was designed to investigate innate immunity elicited by the γ_1 34.5 null mutant upon vaccination. We report that the γ_1 34.5 null mutant induces protective immunity through I κ B kinase in DCs. After adoptive transfer, DCs primed with the γ_1 34.5 null mutant mediate resistance to lethal challenge. This parallels with NF- κ B activation and DC maturation. Notably, the γ_1 34.5 null mutant exerts its activity more effectively in CD8⁺ DCs than in CD8⁻ DCs. Inhibition of NF- κ B phosphorylation or nuclear translocation abrogates the protective effect. Thus, the interaction of the γ_1 34.5 null mutant with CD8⁺ DCs activates NF- κ B, which induces protective immunity.

MATERIALS AND METHODS

Mice. BALB/c mice were purchased from Harlan Sprague-Dawley Inc. and housed under specific-pathogen-free conditions in biosafety level 2 containment. Groups of 5-week-old mice were selected for this study.

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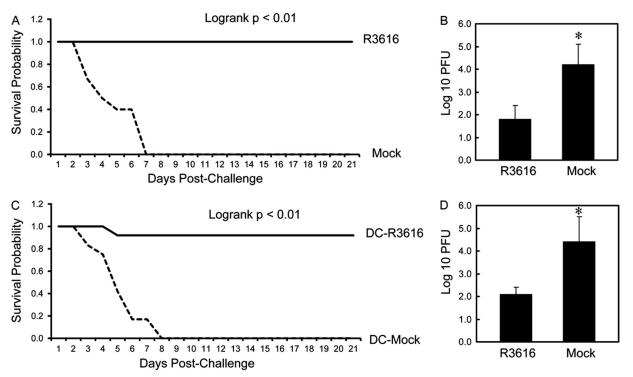


FIG 1 (A) Vaccination with the γ_1 34.5 null mutant induces immunity to HSV-1. Mice were mock inoculated or inoculated with R3616 (1 × 10⁵ PFU) intraperitoneally. Two weeks after immunization, the mice were challenged with HSV-1(F) (1 × 10⁷ PFU) intranasally and monitored over a 21-day period. The survival rates were analyzed by Kaplan-Meier plots (n = 24, log rank P < 0.01). (B) Replication of HSV-1(F). Viral yields in the brain were determined on day 5 after challenge with HSV-1(F) (n = 3 mock-vaccinated control mice, 3 vaccinated mice, P < 0.05). (C) The γ_1 34.5 null mutant-induced protection requires dendritic cells. Mice were mock inoculated or inoculated with R3616 as described for panel A. CD11c⁺ DCs were isolated from spleen as described in Materials and Methods and were transferred into naïve mice on days 1, 3, and 5. On day 6 the mice were challenged with HSV-1(F) (1×10^7 PFU) and monitored for an additional 21 days. The survival rates were analyzed by Kaplan-Meier plots (n = 24, log rank P < 0.01). (D) Replication of HSV-1(F). Viral yields in the brain were determined on day 5 after challenge with HSV-1(F) (n = 3 mock-vaccinated control mice, 3 vaccinated mice, P < 0.05).

Experiments were performed in accordance with the guidelines of the University of Illinois at Chicago.

Cells and viruses. Myeloid DCs were generated as previously described (23). Briefly, bone marrow cells were removed from the tibia and femur bones of BALB/c mice. Following red blood cell lysis and washing, progenitor cells were plated in RPMI 1640 medium (Invitrogen, Auckland, New Zealand) supplemented with 10% fetal bovine serum (FBS), 0.1 mM nonessential amino acids, 1 mM sodium pyruvate, and 20 ng/ml granulocyte-macrophage colony-stimulating factor (GM-CSF; Biosource, Camarillo, CA) in 6-well plates at 4×10^{6} /well. Cells were supplemented with fresh medium every other day. On day 8, DCs were positively selected for surface CD11c expression using magnetic beads (Miltenyi Biotech, Auburn, CA) to give a >97% pure population of CD11c⁺ major histocompatibility complex class II-positive (MHC-II+) cells. DCs displayed low levels of CD40, CD80, CD86, and MHC class II molecules, which is characteristic of immature DCs. Purified CD11c⁺ DCs were cultured in fresh medium with FBS and GM-CSF and used in subsequent experiments. HSV-1(F) is a prototype HSV-1 strain used in this study (14). In recombinant virus R3616, a 1-kb fragment from the coding region of the γ_1 34.5 gene was deleted (8).

Viral infection and DC transfer. Purified CD11c⁺ DCs were plated in 12-well plates (5×10^5 cells/well) or in 96-well round-bottom plates (5×10^4 cells/well) and infected with R3616 (1 or 5 PFU/cell). After 2 h of incubation, cells were washed with phosphate-buffered saline (PBS) and resuspended in RPMI 1640 supplemented with 10% FBS and 20 ng/ml GM-CSF. At different time points after infection, cells were harvested for analysis. For challenge analysis, mice were anesthetized and inoculated intraperitoneally with 1×10^5 PFU of R3616. Two weeks after virus inoculation, mice were intranasally challenged with 1×10^7 PFU of wild-type HSV-1(F) in 50 μ l of PBS. Mice were monitored daily for overall health and sacrificed when symptoms of encephalitis appeared. For in vivo transfer analysis, single splenocyte suspensions were prepared. And CD11c+ DCs (2.0 \times 10⁶ cells/spleen) were isolated and purified by using the CD11c magnetic beads according to the manufacturer's protocol (Miltenyi Biotech). The cells were labeled with fluorescein isothiocyanate (FITC) anti-CD11c⁺ and phycoerythrin (PE) anti-CD8 before sorting. CD11c⁺ CD8⁺ and CD11c⁺ CD8⁻ cells were collected. The cells, with a purity of 96 to 98%, were transferred into naïve mice (5 \times 10⁶ cells/ mouse) three times intraperitoneally. This was done on days 1, 3, and 5, respectively. On day 6 after the first transfer, the mice were challenged with HSV-1(F) and monitored for 3 weeks. For ex vivo analysis, CD11c+ DCs, CD8⁺ DCs, or CD8⁻ DCs derived from bone marrow were mock infected or infected with R3616. At 12 h after infection, the cells were washed with PBS and then adoptively transferred into naïve mice (5 imes10⁶/mouse) intraperitoneally. Mice were then subjected to challenge with wild-type HSV-1(F). For NF-κB inhibition assays, DCs were infected as described above in the presence of a peptide inhibitor, SN50 (50 μ g/ml), or its control peptide, SN50m (Calbiochem, San Diego, CA) (30). Cells were washed and harvested for in vivo or in vitro analysis.

Plaque assay. To determine the titer of infectious virus, on day 5 after challenge with wild-type virus, the brain stems were collected from control and R3616-vaccinated groups (3 to 5 mice) and mechanically homogenized. Samples were serially diluted in 199v medium. Viral yields were titrated on Vero cells and normalized by tissue weight (number of PFU/mg).

Immunoblot analysis and ELISA. To analyze protein expression, cells were harvested and solubilized in disruption buffer containing 50 mM Tris-HCl (pH 7.0), 5% 2-mercaptoethanol, 2% sodium dodecyl sulfate,

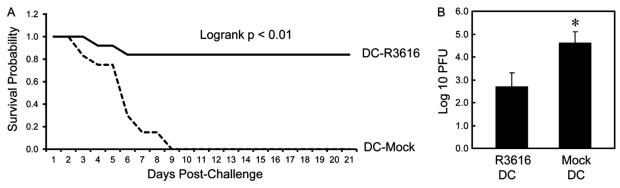


FIG 2 (A) DCs pulsed with the γ_1 34.5 null mutant confer immunity to HSV-1(F). Immature CD11c⁺ DCs from bone marrow were mock infected or infected with R3616 at a multiplicity of infection (MOI) of 5. Cells were washed with PBS at 12 h after infection and adoptively transferred into naïve mice on days 1, 3, and 5. On day 6, mice were challenged with HSV-1(F) (1 × 10⁷ PFU) and monitored over a 21-day period. The survival rates were analyzed by Kaplan-Meier plots (n = 24, log rank P < 0.01). (B) Replication of HSV-1(F). Viral yields in the brain were determined on day 5 after challenge with HSV-1(F) (n = 3 mock-vaccinated control mice, 3 vaccinated mice, P < 0.05).

and 2.75% sucrose. Samples were then subjected to electrophoresis, transferred to nitrocellulose membranes, and reacted with an antibody against FLAG (Sigma), 1 κ B kinase β (IKK β) and phosphorylated IKK β (p-IKK β) (Santa Cruz Biotech, CA), p65/RelA and phosphorylated p65/RelA (p-p65/RelA) (Santa Cruz Biotech), or β -actin (Sigma). The membranes were reacted with either goat anti-rabbit or goat anti-mouse antibody conjugated to horseradish peroxidase (Amersham Pharmacia Biotech, Inc.). Supernatants of cell culture were collected, and IL-12 were quantified using enzyme-linked immunosorbent assay (ELISA) kits from R&D systems according to the manufacturer's instructions.

Cell fractionation assays. Cells were lysed in phosphate-buffered saline containing 0.4% Nonidet P-40 and protease inhibitor mixture (Sigma) and kept on ice with gentle inversion. After brief centrifugation, the nuclei were pelleted and supernatants were collected. After washing, the nuclei were resuspended in phosphate-buffered saline containing 0.4% Nonidet P-40. The cytoplasmic and nuclear fractions were subjected to Western blot analysis with antibodies against p65/RelA (Santa Cruz Biotechnology) and GRP78 (glucose-regulated protein 78; BD Transduction Laboratories) and with antibodies against histone H3 (Cell Signaling), respectively.

Flow cytometry. Tests for cell surface markers CD11c, MHC-II, CD80, and CD86 on DCs were performed according to a standard protocol, with some modifications (23). Cells were blocked with 1 μ l of Fc γ monoclonal antibody (MAb; 0.5 μ g/ml) for 30 min at 4°C. After a onetime wash with PBS, nonpermeabilized cells were stained with isotypematched antibodies, anti-CD11c-PE, anti-MHC-II–FITC, anti-CD80-FITC, and anti-CD86-FITC antibodies, for 30 min on ice with gentle shaking. All antibodies were purchased from eBioscience (San Diego, CA). Samples were processed and screened using a FACSCalibur fluorescence-activated cell sorter (FACS), and data were analyzed with Cell Questpro software (BD).

To determine viral infectivity, CD8⁺ DCs and CD8⁻ DCs mock infected or infected with R3616 were processed as described previously (23). Cells were

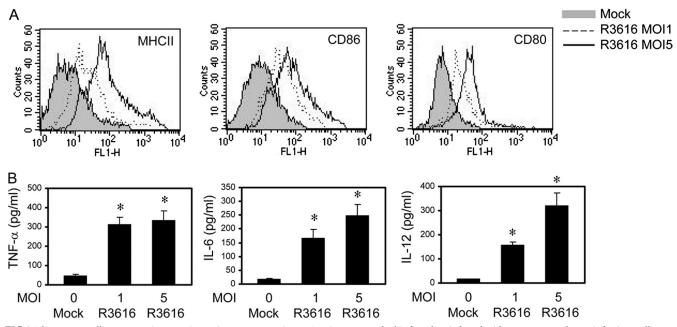


FIG 3 The γ_1 34.5 null mutant activates DCs *ex vivo*. Immature CD11c⁺ DCs were mocked infected or infected with R3616. At 12 h postinfection, cells were assayed for MHC-II, CD86, and CD80 expression by flow cytometry (A). Cell supernatants were assayed for expression of cytokines TNF- α , IL-6, and IL-12 (B). The data are from triplicate samples with standard deviations (*P* < 0.05).

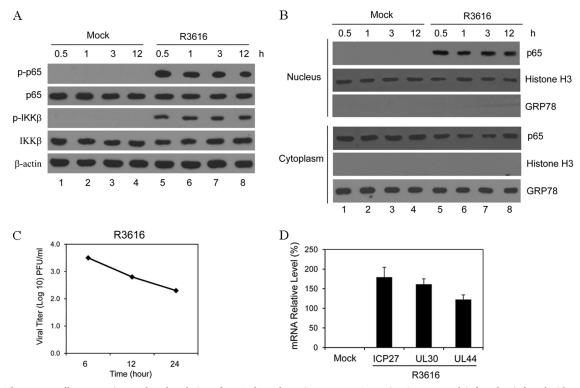


FIG 4 (A) The γ_1 34.5 null mutant triggers phosphorylation of p65/RelA and IKK β . Immature CD11c⁺ DCs were mock infected or infected with R3616 (5 PFU per cell). At the indicated time points, lysates of cells were processed for Western blot (WB) analysis with antibodies against p65/RelA, phosphorylated p65/RelA, IKK β , phosphorylated IKK β , and β -actin. (B) The γ_1 34.5 null mutant stimulates p65/RelA nuclear translocation in DCs. Cells were infected as described for panel A, and the cytoplasmic and nuclear fractions were prepared for Western blot analysis with antibodies against p65, GRP78, and histone H3. (C) Growth of R3616 in DCs. Immature CD11c⁺ DCs were infected with R3616 as described above, and viral titers were measured at the indicated time points. (D) Viral gene expression. Immature CD11c⁺ DCs were infected with R3616, and the expression of ICP27, UL30, and UL44 was determined by quantitative real-time RT-PCR. The results are from triplicate samples with standard deviations.

treated with permeabilizing buffer (eBioscience, San Diego, CA), incubated with a MAb against HSV-1 ICP27 (Virusys, Sykesville, MD), and then reacted with a goat anti-mouse FITC-conjugated antibody (Santa Cruz Biotech, CA). ICP27 expression was evaluated by flow cytometry.

Quantitative real-time PCR. Total RNAs from mock-infected or virusinfected DCs were used to synthesize cDNA as suggested by the manufacturer (Invitrogen). Real-time PCR analysis was performed for ICP27, UL30, and UL44 with the SYBR green system, and all data are presented as relative expression units after normalization to 18S rRNA (24). **Statistics analysis.** The statistical significance of survival rates was determined by the log rank test and Kaplan-Meier survival analysis. Viral titer data, cytokine production, and cell surface marker and RNA expression were evaluated by Student's *t* test. A *P* value of <0.05 was considered statistically significant.

RESULTS

Immunization with the γ_1 34.5 null mutant confers resistance to wild-type HSV-1 through dendritic cells. To establish an *in vivo*

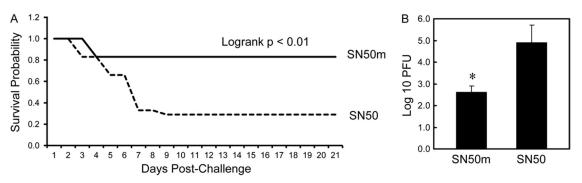


FIG 5 (A) Inhibition of NF- κ B attenuates the γ_1 34.5 null mutant-induced protective immunity. Immature CD11c⁺ DCs were infected with R3616 in the presence of a peptide inhibitor of NF- κ B, SN50, or a control peptide, SN50m (30). At 12 h after infection, cells were transferred into naïve mice on days 1, 3, and 5. On day 6, mice were challenged with HSV-1(F) (1 × 10⁷ PFU) and monitored over a 21-day period. The survival rates were analyzed by Kaplan-Meier plots (*n*) = 24, log rank *P* < 0.01). (B) Replication of HSV-1(F). Viral yields in the brain were determined on day 5 after challenge with HSV-1(F) (*n* = 5 mock-vaccinated control mice, 5 vaccinated mice (*P* < 0.05).

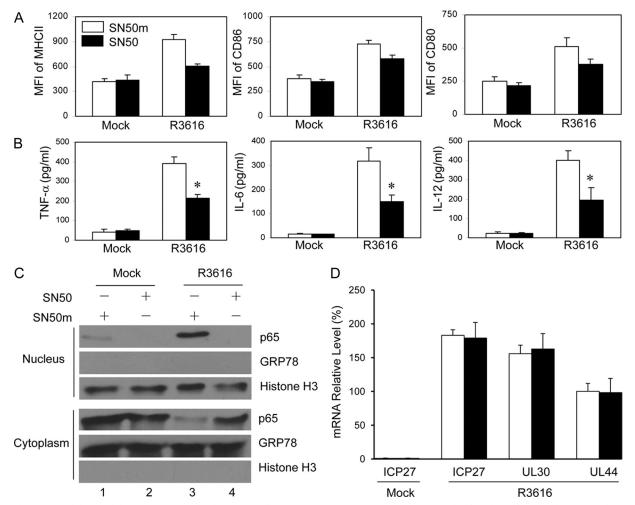


FIG 6 The NF- κ B inhibitor inhibits DC activation by the γ_1 34.5 null mutant. Immature CD11c⁺ DCs were mock infected or infected with R3616 in the presence of SN50 or SN50m. At 12 h postinfection, cells were stained for the expression of MHC-II, CD86, and CD80 by flow cytometry (A). MFI, median fluorescence intensity. Cell supernatants were assayed for TNF- α , IL-6, and IL-12 (B). *, *P* < 0.05. The cytoplasmic and nuclear fractions were prepared and samples were subjected to Western blot analysis with antibodies against p65/RelA, GRP78, and histone H3 (C). The data are representative of results from two independent experiments. (D) Effect of NF- κ B inhibitor on viral gene expression. DCs were mock infected or infected with R3616 as described for panel A. The expression of ICP27, UL30, and UL44 was determined by quantitative real-time PCR, with standard deviations among triplicate samples.

experimental model, we initially assessed the capacity of the γ_1 34.5 null mutant to induce protective immunity in BALB/c mice. Similar to a previous observation (41), vaccination with the γ_1 34.5 null mutant R3616 protected mice against lethal doses of wild-type HSV-1(F) and reduced viral replication in the brain (Fig. 1A and B). Compared to the mock-vaccinated control, no vaccinated mice displayed signs of encephalitis over a 21-day period. To explore the nature of observed phenotypes, we focused on DCs, which are sentinels of host immunity. As such, CD11c⁺ DCs isolated from R3616-vaccinated or mock-vaccinated mice were transferred into naïve mice on days 1, 3, and 5. Mice were challenged with wild-type HSV-1(F) intranasally on day 6 and monitored for 3 weeks. As indicated in Fig. 1C, most recipients of DCs from R3616-vaccinated mice were resistant to wild-type virus. Among mice examined, 91.6% survived. In sharp contrast, none of the recipients of DCs from mock-immunized mice survived after day 8 (Fig. 1C). Consistently, adoptive transfer of DCs from R3616-immunized mice drastically suppressed HSV-1(F) spread to the brain, with the titer being 1.1×10^2 PFU (Fig. 1D). On the

other hand, the control mice failed to suppress HSV-1(F) infection, with the titer increasing to 2.5×10^4 PFU (Fig. 1D). Thus, DCs from mice vaccinated with the γ_1 34.5 null mutant protect against lethal infection from wild-type virus.

Dendritic cells primed with the γ_1 34.5 null mutant mediate protection that parallels with NF- κ B activation. DCs recognize distinct components of HSV (22, 48, 50). When pulsed with a gD peptide *ex vivo*, DCs confer protective immunity upon adoptive transfer (58). To assess the γ_1 34.5 null mutant, immature CD11c⁺ DCs derived from bone marrow were mock infected or infected with R3616 *ex vivo* for 12 h. After washing, the DCs were transferred into naïve mice and challenged with wild-type virus (1 × 10⁷ PFU). As indicated in Fig. 2A, 83% of mice that received R3616-infected DCs were resistant to lethal challenge with HSV-1(F), whereas none of the mice harboring mock-infected DCs survived. In correlation, spread of HSV-1(F) into the brain was significantly suppressed after adoptive transfer of R3616-infected DCs compared to mock-infected DCs (Fig. 2B). These results suggest that DCs primed by the γ_1 34.5 null mutant *ex vivo* initiate protective immunity.

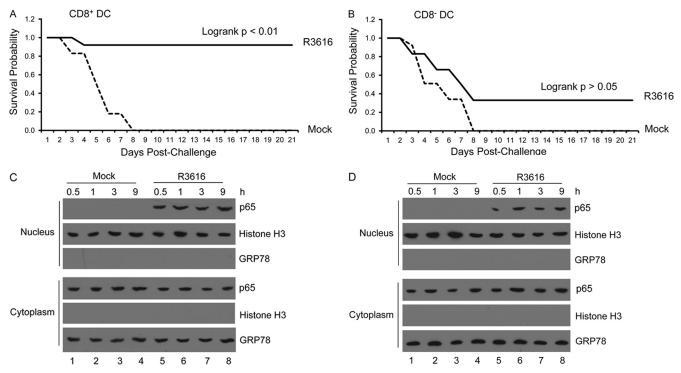


FIG 7 Effects of DC subsets on the $\gamma_1 34.5$ null mutant-induced protective immunity. Immature CD11c⁺ CD8⁺ and CD11c⁺ CD8⁻ DCs were isolated as described in Materials and Methods. CD11c⁺ CD8⁺ (A) or CD11c⁺ CD8⁻ (B) DCs were mocked infected or infected with R3616 (5 PFU per cell). At 12 h after infection, cells were transferred into naïve mice on days 1, 3, and 5. On day 6, mice (n = 24) were challenged with HSV-1(F) (1×10^7 PFU) and monitored over a 21-day period. The survival rates were analyzed by Kaplan-Meier plots. The $\gamma_1 34.5$ null mutant stimulates p65/RelA nuclear translocation in CD11c⁺ CD8⁺ (C) and CD11c⁺ CD8⁻ (D) DCs. Cells were infected as described above, and the cytoplasmic and nuclear fractions were subjected to Western blot analysis with antibodies against p65 and GRP78 and with antibodies against histone H3, respectively. The data are representative of results from three independent experiments.

To examine the basis of the R3616-induced effect, we analyzed DC maturation, which is required for protective immunity (11). Unlike immature DCs, mature DCs have high expression of MHC class II, CD86, CD80, and inflammatory cytokines. Therefore, immature CD11c+ DCs, either mock infected or infected with R3616, were assayed for the expression of these molecules. As shown in Fig. 3A, basal levels of MHC class II, CD86, and CD80 were seen in mock-infected DCs. Infection of immature DCs with R3616 dramatically enhanced the expression of these costimulatory molecules. Moreover, R3616 stimulated the production of cytokines in a dose-dependent manner. As illustrated in Fig. 3B, TNF- α expression remained at basal levels (46 pg/ml) in mockinfected cells. However, its expression increased sharply (>300 pg/ml) after infection with R3616. A similar expression profile was seen for IL-6 and IL-12. Thus, the γ_1 34.5 null mutant stimulates DC maturation.

As DC maturation is coupled to NF- κ B activation (49), we further assessed whether the γ_1 34.5 null mutant had an impact on the NF- κ B signaling pathway. As illustrated in Fig. 4A, while constitutively expressed, both IKK β and p65/RelA remained unphosphorylated in mock-infected DCs. R3616 induced rapid phosphorylation of these proteins after infection. Consistently, p65/ RelA was primarily located in the cytoplasm fraction in mockinfected cells (Fig. 4B). After R3616 infection, p65/RelA appeared in the nuclear fraction at all time points tested. R3616 infection had no effect on control proteins histone 3 in the nucleus and GRP78 in the cytoplasm, suggesting that R3616 infection activated I κ B kinase and NF- κ B. While unable to produce infectious virus, R3616 expressed ICP27, UL30, and UL44 mRNA in infected DCs (Fig. 4C and D). We conclude that the γ_1 34.5 null mutant abortively infects DCs and activates NF- κ B, which coincides with the induction of costimulatory molecules, cytokines, and protective immunity.

Inhibition of NF-kB activation in dendritic cells disrupts protective immunity elicited by the γ_1 34.5 null mutant. Based on the above-described analyses, we reasoned that the γ_1 34.5 null mutant-induced NF-KB activation in DCs might be a determinant of protective immunity. To test this hypothesis, we carried out the NF-kB inhibition-coupled adoptive transfer assays. Immature CD11c⁺ DCs were infected with R3616 in the presence of a cellpermeant peptide NF-*k*B inhibitor, SN50, or its control peptide, SN50m (30). At 12 h after infection, cells were washed and adoptively transferred into naïve mice for challenge analysis. As illustrated in Fig. 5A, the majority of mice that received DCs with the NF- κ B inhibitor did not survive after challenge with wild-type HSV-1(F). In this treatment group, only 25% of the animals were protected. However, most mice that received DCs with a control inhibitor survived HSV-1(F) challenge. Eighty-three percent of animals in this treatment group were protected. Concordant with these results, the NF- κ B inhibitor SN50 enhanced viral spread to the brain after HSV-1(F) infection compared to the control inhibitor SN50m, with the titer reaching 8×10^4 PFU on day 5 (Fig. 5B). Thus, the γ_1 34.5 null mutant-induced protective immunity is impaired by the NF- κ B inhibitor.

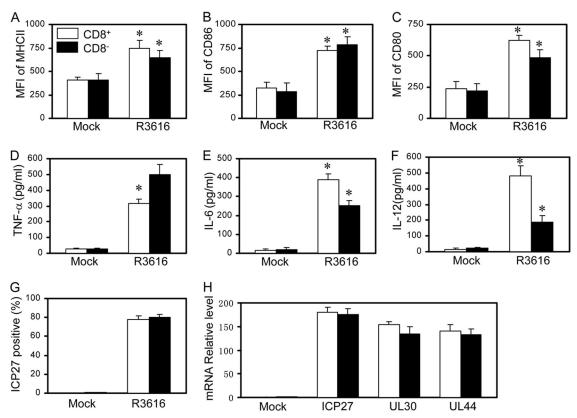


FIG 8 The γ_1 34.5 null mutant triggers DC subsets differentially. Immature CD11c⁺ CD8⁺ and CD11c⁺ CD8⁻ DCs were mocked infected or infected with R3616 (5 PFU per cell). At 12 h after infection, cells were assayed for MHC-II (A), CD86 (B), and CD80 (C) expression by flow cytometry. Cell supernatants were assayed for expression of cytokines TNF- α (D), IL-6 (E), and IL-12 (F). (G) Viral infectivity was determined by examining ICP27 expression as described in Materials and Methods. (H) The expression of ICP27, UL30, and UL44 in DCs infected with R3616 was determined by quantitative real-time PCR. The results are from triplicate samples with standard deviations (*, *P* < 0.05).

To further evaluate the effect of the NF- κ B inhibitor, we assessed DC maturation by R3616 ex vivo. As shown in Fig. 6A and B, unlike mock infection, R3616 infection stimulated the expression of MHC-II, CD80, CD86, TNF- α , IL-6, and IL-12 in the presence of the control, SN50m. Addition of SN50 resulted in lower levels of costimulatory molecules and cytokines. It seems that the inhibitory effect by the NF-kB inhibitor was more drastic on the production of cytokines than on costimulatory molecule expression. Notably, the NF-KB inhibitor SN50 but not the control, SN50m, effectively blocked nuclear translocation of p65/RelA (Fig. 6C). Nevertheless, treatment with SN50 or SN50m had no effect on viral gene expression where R3616 expressed ICP27, UL30, and UL44 genes (Fig. 6D), suggesting that SN50 directly inhibited NF-KB activation. These results argue that NF-KB activation in DCs by the γ_1 34.5 null mutant is essential to induce protective immunity against HSV-1 infection in vivo.

The γ_1 34.5 null mutant exerts differential effects through CD8⁺ and CD8⁻ dendritic cells. Conventional DC subsets can generally be categorized as CD8⁺ DCs and CD8⁻ DCs. To investigate how the γ_1 34.5 null mutant exerted its effect, we analyzed CD8⁺ DCs and CD8⁻ DCs derived from bone marrow (>98% purity). Cells were either mock infected or infected with R3616 for 12 h. After washing, cells were transferred into naïve mice for challenge assays. As shown in Fig. 7A, 91.6% of mice that received R3616-infected CD8⁺ DCs survived after HSV-1(F) challenge. As expected, none of the mice that received mock-infected CD8⁺

DCs were viable beyond day 8. Interestingly, only 33.3% of mice that received R3616-infected CD8⁻ DCs survived lethal challenge (Fig. 7B). A large fraction of mice within this treatment group did not survive beyond day 8. This lack of protection resembled the phenotype seen in mice receiving mock-infected CD8⁻ DCs. Hence, CD8⁺ DCs confer protection more effectively than CD8⁻ DCs after priming by the γ_1 34.5 null mutant *ex vivo*.

To understand the observed differences, we examined the status of NF-KB activation ex vivo. As shown in Fig. 7C and D, R3616 induced nuclear translocation of p65/RelA both in CD8+ DCs and in CD8⁻ DCs. Next, we evaluated DC maturation. As illustrated in Fig. 8A to C, in response to R3616 infection, CD8⁺ DCs as well as CD8- DCs exhibited enhanced expression of MHC II, CD80, and CD86 at comparable levels. Nonetheless, CD8⁺ DCs secreted a higher level of IL-6 and IL-12, whereas CD8⁻ DCs expressed more TNF- α (Fig. 8D to F). Under these conditions, R3616 infected CD8⁺ DCs and CD8⁻ DCs equally well, with approximately 80% infectivity, as revealed by FACS analysis of ICP27 (Fig. 8G). As infection continued, R3616 expressed comparable levels of ICP27, UL30, and UL44 in CD8⁺ DCs and CD8⁻ DCs, as measured by quantitative reverse transcription-PCR (RT-PCR) analysis (Fig. 8H). Therefore, although it activates NF- κ B in CD8⁺ DCs and CD8⁻ DCs, the γ_1 34.5 null mutant induces cytokine expression differentially.

The γ_1 34.5 null mutant-induced immunity requires NF- κ B activation in CD8⁺ dendritic cells. Finally, we tested whether the

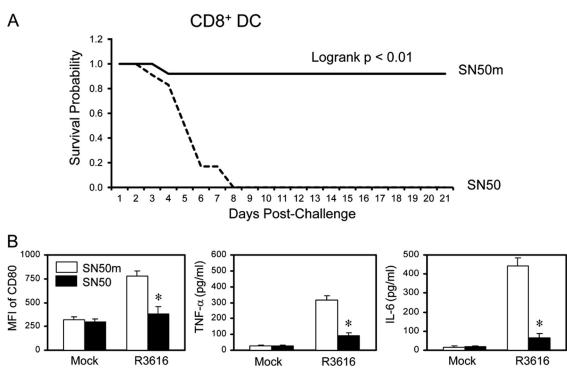


FIG 9 (A) CD11c⁺ CD8⁺ DC-induced immunity requires NF-κB activation. Immature CD11c⁺ CD8⁺ DCs were infected with R3616 in the presence of SN50 or SN50m (30). At 12 h after infection, cells were transferred into naïve mice on days 1, 3, and 5. On day 6, mice were challenged with HSV-1(F) (1 × 10⁷ PFU) and monitored over a 21-day period. The survival rates were analyzed by Kaplan-Meier plots (n = 24, log rank P < 0.01). (B) Immature CD11c⁺ CD8⁺ DCs infected as described for panel A were assayed for CD80, TNF- α , and IL-6 expression. The results are from triplicate samples with standard deviations (*, P < 0.05).

 γ_1 34.5 null mutant exerted its effect through NF-κB in CD8⁺ DCs. Purified CD8⁺ DCs were infected with R3616 in the presence of SN50 or SN50m for 12 h. Cells were adoptively transferred into naïve mice, which were challenged with HSV-1(F). As illustrated in Fig. 9A, none of the mice that received DCs with the NF-κB inhibitor SN50 survived beyond day 8 after lethal challenge with HSV-1(F). In stark contrast, 92% of mice that received DCs with the control inhibitor SN50m survived. In correlation, SN50 but not SN50m effectively inhibited the induction of CD80, TNF- α , and IL-6 by R3616 in CD8⁺ DCs (Fig. 9B). These results suggest that the γ_1 34.5 null mutant-induced protective immunity requires NF-κB activation in CD8⁺ dendritic cells.

DISCUSSION

Control of acute or persistent HSV infection initially involves the innate immune system, which consists of DCs, macrophages, natural killer cells, cytokines, and complement proteins (11). In this study, we show that DCs from mice vaccinated with an HSV mutant lacking the $\gamma_1 34.5$ gene confer resistance to HSV-1. *Ex vivo* exposure of DCs to the $\gamma_1 34.5$ null mutant activates NF- κ B, which parallels with upregulation of costimulatory molecules and inflammatory cytokines. This interaction translates into protective immunity against wild-type HSV-1 *in vivo*. Inhibition of NF- κ B signaling in DCs reverses the protective effect. CD8⁺ DCs function more effectively than CD8⁻ DCs. These results support the concept that stimulation of innate immune signaling by the $\gamma_1 34.5$ null mutant potentiates protective immunity.

The γ_1 34.5 null mutant is avirulent *in vivo* (4, 8, 33, 35, 43, 51, 56, 63). Although unable to produce infectious virus in DCs, the

 γ_1 34.5 null mutant expressed early and late genes. Upon adoptive transfer, these DCs mediated protection against lethal challenge from wild-type virus. A question arises as to how the γ_1 34.5 null mutant elicits immunity via DCs. Accumulating evidence suggests that TLR3, MDA5, and RIG-I detect HSV RNA (36, 47, 67), whereas polymerase III and IFI16 recognize intracellular HSV DNA (7, 59). Additionally, an unknown receptor in DCs senses HSV glycoproteins (50). We speculate that viral DNA, RNA, or a protein from the γ_1 34.5 null mutant may trigger an aforementioned pathway(s) in its abortive infection, leading to protective immunity. In this respect, it is notable that HSV mutants deficient in ICP0, ICP4, or ICP27 behave differently in DCs (37). The ICP27 mutant stimulates the expression of type I interferon (IFN) and inflammatory cytokines in DCs, whereas ICP0 and ICP4 mutants do not. This is thought to result from an effect of ICP27 on posttranscriptional events in host RNA processing or the expression of a viral inhibitor(s) of cytokine expression (37). DCs recognize HSV via TLR-dependent and -independent pathways (22, 48, 50). As ICP27 and γ_1 34.5 mutants differ in the nature of mutation, such a difference may affect their ways to interact with DCs. Although activating DCs, these mutants likely trigger innate immune pathways differently. Additional work is needed to test this hypothesis.

I κ B kinase sits at the center of innate immune pathways, and its activation is linked to DC maturation (27, 49). It is well recognized that HSV both activates and inhibits NF- κ B during infection. While NF- κ B activation by HSV is required for optimal viral replication and cell survival in some cell settings (16, 17, 42), its activation also stimulates antiviral immunity (12, 37, 60). Intrigu-

ingly, HSV activation of NF-κB is linked to double-strandeddependent protein kinase PKR (57). We noted that the γ_1 34.5 null mutant induced phosphorylation of IKK β and p65/RelA in DCs which induced protective immunity upon adoptive transfer. Inhibition of NF-κB activation reversed these phenotypes but had no effect on viral gene expression. We suspect that upon infection with the γ_1 34.5 null mutant, viral proteins, such as gD and UL37, may activate NF-κB (31, 53). Alternatively, viral nucleic acids may stimulate NF-κB activation in DCs. Although not investigated, our data do not exclude the idea that the γ_1 34.5 null mutant may act on DCs through additional pathways. An attractive possibility is to activate interferon regulatory factor 3 (IRF3), which leads to the expression of type I IFN and chemokines (27, 61). Another possibility is to stimulate transcription factor AP-1, which controls cytokine expression (27, 66).

It is noteworthy that the γ_1 34.5 null mutant affects CD8⁺ DCs and CD8⁻ DCs differentially. Although activating NF-*k*B both in CD8⁺ DCs and in CD8⁻ DCs, the γ_1 34.5 null mutant induced protective immunity only via CD8⁺ DCs. We noted that the γ_1 34.5 null mutant infected CD8⁺ DCs and CD8⁻ DCs equally well, with comparable early and late gene expression. This result rules out a difference in viral infection that may contribute to the observed phenotypes. A plausible explanation is that these DC subsets are functionally distinct. Therefore, besides NF-KB activation, an additional signal(s) or component from DCs is required to initiate protective immunity. Interpreted within this model, it is interesting that CD8⁺ DCs secreted a higher level of IL-6 and IL-12 than CD8⁻ DCs upon exposure to the γ_1 34.5 null mutant. Conversely, CD8⁻ DCs produced more TNF- α . As these DC subsets displayed similar levels of cell surface molecules, different cytokine responses probably contributed to the phenotypes observed in vivo.

CD8⁺ DCs have been reported to play a key role in controlling viral infections (1, 2, 54). In light of these observations, it is intriguing that the γ_1 34.5 null mutant induced protective immunity through CD8⁺ DCs. In principle, the γ_1 34.5 null mutant may primarily target CD8⁺ DCs upon immunization. In this process, it likely promotes DC maturation as well as antigen presentation, which is coupled to NF- κ B activation (64, 65). Emerging evidence suggests that CD8⁺ DCs preferentially initiate CD8 T cell immunity through TLR3 in response to HSV infection (13). Hence, the γ_1 34.5 null mutant may activate the TLR3 pathway leading NF- κ B activation. In line with this model, the γ_1 34.5 null mutant induced NF-*k*B activation and the maturation of CD8⁺ DCs. While having no effect on viral gene expression, inhibition of NF-kB disrupted the ability of CD8⁺ DCs to mediate protective immunity in vivo. These observations underscore a key role of NF-KB and CD8+ DCs in controlling HSV infection. Work is in progress to investigate the underlying mechanisms.

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