



Herpesvirus Entry Mediator and Ocular Herpesvirus Infection: More than Meets the Eye

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ABSTRACT As its name suggests, the host receptor herpesvirus entry mediator (HVEM) facilitates herpes simplex virus (HSV) entry through interactions with a viral envelope glycoprotein. HVEM also bridges several signaling networks, binding ligands from both tumor necrosis factor (TNF) and immunoglobulin (Ig) superfamilies with diverse, and often opposing, outcomes. While HVEM was first identified as a viral entry receptor for HSV, it is only recently that HVEM has emerged as an important host factor in immunopathogenesis of ocular HSV type 1 (HSV-1) infection. Surprisingly, HVEM exacerbates disease development in the eye independently of entry. HVEM signaling has been shown to play a variety of roles in modulating immune responses to HSV and other pathogens, and there is increasing evidence that these effects are responsible for HVEM-mediated pathogenesis in the eye. Here, we review the dual branches of HVEM function during HSV infection: entry and immunomodulation. HVEM is broadly expressed; intersects two important immunologic signaling networks; and impacts autoimmunity, infection, and inflammation. We hope that by understanding the complex range of effects mediated by this receptor, we can offer insights applicable to a wide variety of disease states.

KEYWORDS HVEM, herpes simplex virus, herpes stromal keratitis

Herpes simplex virus 1 (HSV-1) infects the majority of the world's population by adulthood and is responsible for the vast majority of ocular herpesvirus infections (1–3). In humans and mice, HSV-1 establishes lifelong latency in the trigeminal ganglia (TG) (4, 5). Reactivations of HSV-1 in the TG can lead to anterograde movement of the virus along the ophthalmic branch of the trigeminal nerve, resulting in recurrent infection of virtually all the superficial tissues of the eye, including the cornea, conjunctiva, and eyelid (6–8). Ocular herpesvirus infections can lead to epithelial ulceration of the cornea, uveitis, and retinitis but most commonly cause herpes stromal keratitis, or HSK (9). HSK is characterized by chronic inflammation of the corneal stroma, leading to corneal thickening, opacification, scarring, and, potentially, blindness (10, 11). While reactivation accounts for the majority of human disease, most murine studies of HSK model primary infection as mice do not efficiently or reliably reactivate from latency (6).

In the primary murine model of HSK, actively replicating HSV-1 in the cornea is detectable by plaque assay for up to 5 to 6 days postinfection (dpi) (9, 12). Viral replication initiates HSK, as UV-inactivated or replication-deficient mutant HSV fail to induce HSK in BALB/c mice (13). However, HSK is an inflammatory disease, brought about by host infiltrates, particularly CD4⁺ T cells and polymorphonuclear cells (PMN), that invade the murine cornea several days after replicating virus has been cleared and that persist chronically (14–18). A dizzying array of cytokines, chemokines, and immune cell types have been implicated in the pathogenesis of HSK, the temporal and functional relationships of which have been expertly reviewed elsewhere (11, 15, 18–20). Here, we focus on a single host factor, herpesvirus entry mediator (HVEM, also called

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tumor necrosis factor [TNF] superfamily member receptor 14 [*Tnfrsf14*]), originally identified as a viral entry receptor. HVEM has a multitude of roles in mucosal responses to a variety of pathogens (21–25) and has been found to influence HSV pathogenesis in the murine eye during most stages of infection, including entry/acute viral replication, early/innate responses, chronic inflammation, and even viral latency. Results from our laboratory indicate that HVEM functions as a proinflammatory factor in murine herpes keratitis and call into question its importance as an entry receptor during ocular infection. However, the role of HVEM in immunopathogenesis of murine HSK is still emerging, and its importance during human ocular herpesvirus infections remains unexplored.

HVEM IS A VIRAL ENTRY RECEPTOR

Entry mechanics of HSV. Herpesviruses are large, enveloped, double-stranded DNA viruses (26). Glycoproteins studding the viral envelope are essential for entry into host cells, a complex process requiring at least four separate glycoproteins, including glycoprotein B (gB), gH/gL, and gD (27). Fusion between viral and host membranes requires binding of gD to a host surface receptor (27). Several surface proteins have been identified as gD receptor targets, including nectin-2, HVEM, and nectin-1, although nectin-1 is considered the most biologically relevant entry receptor for HSV (28–31). Nectin-1 is involved in cell-cell adhesion and is a member of the immunoglobulin (Ig) superfamily (32), while HVEM is a TNF receptor superfamily member (28).

The crystal structures of gD bound to both nectin-1 and HVEM have been solved (33, 34). Binding of HVEM or nectin-1 occurs on distinct regions of gD dimers (35), although binding by either ligand triggers similar conformational changes in the structure of the viral glycoprotein, exposing the C-terminal pro-fusion domain (36). HSV entry is a complex process that has been reviewed elsewhere (27), but a highly simplified description of the mechanism is as follows: glycoprotein D interacts with gB trimers bound to a gB-specific receptor and to the heterodimer gH-gL, which bridges receptor binding with gB activation and fusion (27, 37). The sum of these interactions is the insertion of the gB fusion loops into the cellular plasma membrane (or, in some circumstances, into the endosomal membrane) and refolding of the gB trimer into a postfusion form (29). Subsequent mixing of viral and host membranes and the formation of a fusion pore through which the viral capsid and tegument proteins can be deposited complete the entry process (38).

While HVEM and nectin-1 exhibit similar kinetics and affinities of binding to gD *in vitro* (37, 39), early *in vivo* studies showed a greater importance for nectin-1 in the pathogenesis of HSV, especially in terms of invasion of and spread throughout the nervous system in intravaginal and intracranial models of HSV-2 infection (40, 41). Remarkably, nectin-1 knockout (KO) mice inoculated with HSV-2 directly into the hippocampus do not develop encephalitis, despite the presence of HVEM in the brain; they also lack demonstrable virus by immunofluorescence (40). Although HVEM is largely dispensable for HSV-2 infection of the brain (40), vagina (41), or eye (42), our laboratory has shown that HVEM promotes HSV-1 pathogenesis specifically in the setting of murine ocular infection (43, 44). In order to understand why this receptor is required for pathogenesis of ocular herpetic infections, both the entry and immunomodulatory functions of HVEM must be explored.

HVEM-mediated entry in the eye. HVEM expression has been evaluated in a number of ocular tissues. Results of studies of cultured human cell lines by real-time PCR (RT-PCR) and immunofluorescence or flow cytometry indicate that retinal pigment epithelial (RPE) cells (45), corneal fibroblasts (46), trabecular meshwork cells (47), and conjunctival and corneal epithelial cells (48, 49) express HVEM mRNA and membrane-bound protein. Use of HVEM-blocking antibodies (47, 49) or HVEM small interfering RNA (siRNA) knockdown (48) reduces viral entry of some, but not all, of these cell types *in vitro*, suggesting that HVEM is the entry receptor in use. However, widespread expression of nectin-1 and the sufficiency of this receptor for *in vivo* infection of the murine cornea and TG preclude the notion that HVEM is the primary receptor in the eye (43,

50–52). Our experiments performed with a well-characterized HSV-1 mutant, HSV-1 (17)gD Δ 7-15, which is restricted to nectin-1 entry through targeted deletion of the HVEM-specific binding region of gD (29, 31, 34, 35, 39, 53, 54), confirmed that HVEM is not the primary entry receptor in the cornea: infection with the gD Δ 7-15 mutant produces titers, clinical disease, and inflammatory cytokine levels equivalent to those seen with the HVEM-competent virus control (44). Furthermore, recent analysis of whole murine corneas by flow cytometry indicated that HVEM expression on epithelial or endothelial cells is limited *in vivo* (55).

This is not to say that HVEM is unimportant during ocular HSV-1 infection. Despite the presence of a suitable alternate entry receptor, nectin-1, in the cornea, HSV-1 infection of *Tnfrsf14*^{-/-} (HVEM KO) mice results in lower viral loads in eye swabs, with subsequent loss of titer in the TG, brain, and periocular skin (POS), and reduced rates of reactivation from the TG (42, 43). Along with this replication defect, HVEM KO mice are protected from systemic clinical symptoms of HSV-1, including lesion development and neurologic morbidity (42, 43), as well as from cornea-specific pathology, including loss of sensitivity to mechanical pressure (55), inflammatory cytokine release, and stromal leukocytic infiltration (44). Entry-independent contributions of HVEM during HSV-1 pathogenesis likely stem from tuning of immune responses by HVEM during inflammation. While we are convinced that HVEM contributes to inflammation and immunopathology during murine HSK, the precise mechanisms underlying this process remain largely unknown. Evidence gleaned from other disease models/organs in which HVEM signaling has been more thoroughly explored provides clues about how HVEM may promote inflammation in the eye.

HVEM ALTERS HOST IMMUNE RESPONSES TO HSV

HVEM signaling in the host. HVEM, also designated CD270, is a bidirectional receptor that can bind ligands of both the TNF and Ig superfamilies (Fig. 1), producing a wide diversity of outcomes (25). Outside of the eye, HVEM expression is broad: although its expression fluctuates throughout maturation, HVEM is found on most types of leukocytes, including T and B cells, dendritic cells (DCs), natural killer (NK) cells, and myeloid cells (56). HVEM is also highly expressed by nervous tissue and gut and lung epithelia (23, 57).

Identified HVEM ligands include LT α (lymphotoxin α), LIGHT (lymphotoxin-related inducible ligand that competes for glycoprotein D binding to HVEM on T cells), BTLA (B and T lymphocyte attenuator), and CD160 (58–60). Immature DCs, monocytes, and activated T cells express membrane-bound or soluble forms of LT α and LIGHT (56). These trimeric TNF family ligands typically enhance activation or differentiation of a variety of immune cell types upon binding cysteine-rich domain 2 (CRD2) and CRD3 of HVEM (61–64). BTLA is an Ig superfamily member found on T and B cells, DCs, and myeloid cells (59). CD160 is a dimeric, glycosylphosphatidylinositol (GPI)-anchored protein with an Ig-like fold that is prototypically expressed by NK cells but is also found on subsets of CD4⁺ and CD8⁺ T cells (60). Both BTLA and CD160 communicate a corepressive signal upon binding HVEM CRD1 in most described cases (65–69), although CD160 is known to be activating in some circumstances, i.e., on NK cells, where it is required for interferon gamma (IFN- γ) production (70). However, the outcomes of HVEM signaling differ depending on whether the ligand is soluble or in the membrane-bound form, whether the interaction occurs in *cis* or in *trans*, and on the specific identity of the cells involved (25, 60, 71, 72). Viral gD also binds CRD1 and most directly competes with BTLA for HVEM binding, although there is evidence indicating that LIGHT and LT α interactions with HVEM may also be affected by the presence of gD (73). Because the TNF- and Ig-ligand domains occur on different faces of the receptor, combinations of HVEM and its ligands may result in ternary complexes (64, 65, 74–76).

As a receptor, HVEM engagement by LIGHT, CD160, BTLA, or viral gD recruits members of the TNF receptor-associated factor (TRAF) family to the cytoplasmic tail of HVEM and activates prosurvival nuclear factor κ B (NF- κ B) signaling (77, 78). NF- κ B is a transcription factor normally sequestered in the cytoplasm by proteins, including I κ B

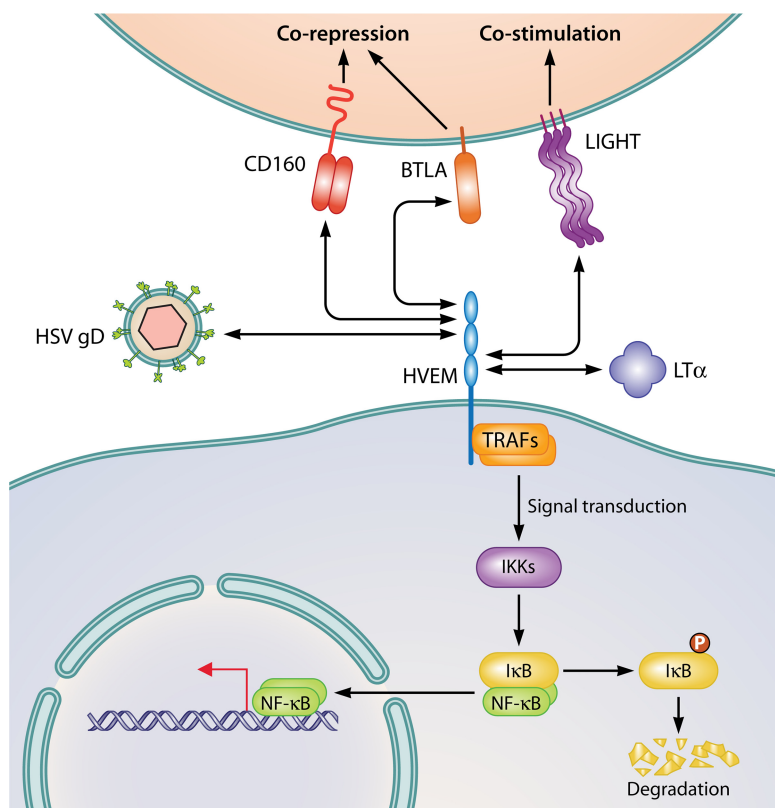


FIG 1 HVEM signaling is bidirectional. HVEM is a TNF receptor superfamily member that can interact with Ig-like ligands (CD160 and BTLA), TNF ligands (LIGHT and LT α), and an HSV glycoprotein, gD. HVEM is expressed on a broad range of cell types, including T cells, B cells, DCs, NK cells, macrophages, PMN, neurons, and epithelial cells. BTLA and LIGHT are also found on most leukocytes, including B and T cells, granulocytes, NK cells, etc. CD160 is found on subsets of CD4⁺ and CD8⁺ T cells, NK cells, and intraepithelial lymphocytes (IELs). BTLA, CD160, and HSV gD bind cysteine-rich domain 1 (CRD1) of HVEM, while LIGHT and soluble LT α bind CRD2 and CRD3. Outcomes can vary from corepressive signals delivered through CD160 and BTLA in *trans* to costimulatory signals delivered through LIGHT. BTLA and LIGHT can also associate with HVEM in *cis*, forming heterotrimeric complexes, the conformation most commonly seen on resting T cells. Binding in *cis* holds HVEM in an inactive state, without NF- κ B activation. In contrast, binding in *trans* of HVEM by any of its ligands leads to the activation of NF- κ B signaling through recruitment of TRAF family members by the cytoplasmic portion of HVEM. Subsequent activation of I κ B kinase/I κ α kinase (IKK β / α), phosphorylation and degradation of the NF- κ B chaperone I κ B, and activation and nuclear translocation of NF- κ B result in increased transcription of inflammatory factors and prosurvival signals within the HVEM-expressing cell.

kinase/I κ α kinase (IKK β / α); upon their degradation, NF- κ B translocates to the nucleus to initiate transcription of DNA, cytokine production, and cell survival (79). HVEM activates prosurvival signaling through NF- κ B even when its binding partner is a corepressive ligand, such as BTLA or CD160. For example, T cells receiving cosuppressive signals through BTLA cease proliferating, while *Btla*^{-/-} T cells stimulated by soluble BTLA-Fc induce NF- κ B signaling via HVEM and exhibit increased survival (78). NF- κ B induced by the binding of gD to HVEM has antiapoptotic effects *in vitro*, and loss of NF- κ B activation (and nuclear translocation), which is sensitive to blocking with anti-gD antibodies, reduces viral yield 80 to 90% and increases apoptosis (80–84). HSV may benefit from HVEM activation through NF- κ B-mediated prosurvival signals that prevent apoptosis of infected cells (80, 83).

Understanding the intricacies of the HVEM signaling network is challenging given the number of possible ligands involved, the lack of predictability with respect to whether an interaction will ultimately be pro- or anti-inflammatory, and the possibility of competing, opposing signals arising from the same receptor-ligand interaction. There is increasing evidence that HVEM modulates aspects of the innate and adaptive

immune responses to HSV in the murine cornea, although the clinical significance of HVEM signaling in human patients with HSK remains to be elucidated.

HVEM and innate immune responses. One of the first indications that herpesviruses may alter early innate immune responses to infection via HVEM came from the murine vaginal model of genital HSV-2 infection. Mice infected with an HVEM-entry null HSV-2 strain (with the $\Delta 7-15$ deletion) have significantly higher levels of the cytokine interleukin-6 (IL-6) and chemokines CXCL9, CXCL10, and CCL4 in vaginal washes than do mice infected by HSV-2 that could engage HVEM (85). Infected HVEM KO corneas have decreased levels of several inflammatory cytokines, including IL-6 and CXCL10, compared to infected C57BL/6 (wild-type [WT]) controls early after infection (44). However, unlike in the vaginal model, the induction of these cytokines in the cornea is independent of HVEM-mediated entry of HSV-1 (44). IL-6 and CXCL10 are neutrophil and T cell chemoattractants, respectively, known to promote ocular HSV-1 pathogenesis (19, 86–89), and increased expression of these chemotactic factors may lead to heavier infiltration by myeloid and lymphoid cells in WT versus HVEM KO murine corneas (44). However, siRNA-mediated knockdown of HVEM from cultured telomere-immortalized human corneal epithelial cells decreased production of IFN- γ , MIP-1 α (CCL3), and MIP-1 β (CCL4) after HSV-1 challenge (90). This discrepancy may represent differences between murine and human cells, or may have arisen because in the *in vitro* system, HVEM stimulation occurred solely with HSV gD, while *in vivo*, binding by natural ligand(s) likely takes precedence (44).

The mechanism by which HVEM upregulates cytokine secretion in the murine cornea has not been defined, although it must involve one of the host HVEM ligands as gD binding is not required (44). Elegant studies of innate responses to bacteria in lung and gut mucosal epithelia have found a similar role for host HVEM signaling in the induction of IL-6 and other cytokines (22). In this model, CD160 on innate-like intraepithelial lymphocytes activates HVEM, which is highly expressed by the intestinal epithelium, resulting in NF- κ B-mediated Stat3 activation and increased expression of genes and peptides related to epithelial immunity (22, 23). During HSK, the corneal epithelium secretes IL-6 and other cytokines in response to HSV-1 infection (91, 92). We performed experiments using adoptive transfer between WT and HVEM KO mice and found that HVEM on a radiation-resistant cell type(s), such as the corneal epithelium, is required for disease progression, while HVEM on a radiation-sensitive cell type(s), such as circulating immune cells, is dispensable (44). However, we were unable to detect substantial quantities of HVEM on the corneal epithelium (or stroma) *in vivo* either by IHC or flow cytometry, making it unlikely that these are the source of the HVEM-mediated cytokine production (55).

Flow cytometry of whole corneas from C57BL/6 mice indicates that the majority of HVEM is located on CD11b⁺ CD11c⁻ Ly6C⁺ Ly6G⁻ monocyte lineage cells in the acute phase and on PMN and CD4⁺ T cells in the chronic phase (55). The shift in HVEM-positive (HVEM⁺) populations likely reflects differences in the cellular composition of the ocular infiltrate over time rather than fluctuations in HVEM expression, although this was not directly tested in our study. Stromal macrophages (and DCs) incompletely turn over with irradiation (93); therefore, these could be the radiation-resistant HVEM⁺ cell type identified as mediating disease in adoptive-transfer experiments (44). Although other HVEM⁺ lineages may also contribute to pathogenesis, albeit in a subtler fashion, monocytes/macrophages are particularly intriguing to us in light of these findings.

Along with NK cells, which play an essential role in the regulation of early HSV replication (94), corneal and conjunctival macrophages are required for viral restriction in the first 48 h after infection (95–98). However, control of viral replication comes at a price: macrophages, along with corneal-infiltrating and lymph node-residing DCs (99), promote CD4⁺ T cell activation during ocular HSV-1 infection (98), although the roles of resident versus recruited macrophages in CD4⁺ T cell activation are poorly defined in HSK. Even without DCs, chronic-phase HSK (beyond day 7) still develops in HSV-

infected mice, potentially through the activation of CD4⁺ T cells by closely associated corneal macrophages (99). Macrophage-associated cytokines such as MIP-1 α and MIP-1 β contribute to further corneal infiltration and damage (19). Macrophages and PMN also promote vascularization of the cornea, a required step in HSK development, through secretion of vascular endothelial growth factor receptor (VEGF) and matrix degradation enzymes (100, 101). All of these effects could be influenced by HVEM signaling on macrophages and neutrophils, potentially through HVEM activation operating directly or through interactions with its ligands.

LIGHT binding of HVEM on macrophages and neutrophils provides an activating signal, increasing phagocytic activity and production of inflammatory/antibacterial factors, including nitric oxide (NO), reactive oxygen species (ROS), IL-8, and TNF- α (102), via changes in intracellular calcium sequestration (103). Agonistic binding of HVEM on neutrophils also increases respiratory burst and degranulation, providing a further explanation for the increased bactericidal activity of PMN via HVEM (104). *In vivo*, secretion of type I interferon from splenic cells, especially macrophages, partially requires HVEM; in HVEM-deficient mice, loss of type I interferons reduces lymphocyte bystander activation and immunopathology after *Listeria* infection (105).

In contrast, HVEM-BLTA interactions reduce activation of innate cell populations, including macrophages, inflammatory monocytes, and PMN (24, 66, 67). During acute experimental sepsis, HVEM-BTLA interactions on innate populations worsen outcomes with respect to organ injury, bacterial burden, and mortality (106). Although both HVEM and BTLA are expressed on recruited myeloid cells in this model, BTLA-directed corepressive signals explain the finding of reduced myeloid activation and survival more convincingly than HVEM-directed prosurvival/NF- κ B signals. Our laboratory is currently investigating the corneal expression of LIGHT and BTLA on resident and infiltrating cells during HSV-1 infection and what role these ligands play in HVEM-mediated disease.

HVEM and adaptive immune responses. Helper Th1 CD4⁺ T cells are considered to represent the major immunopathologic cell type in HSK (9, 10, 15). Investigators have repeatedly shown that without functional CD4⁺ T cells, HSK does not develop (107–109). Corneal CD4⁺ T cells remain activated in the absence of replicating virus, likely through bystander activation (110), in which CD4⁺ T cells become nonspecifically activated due to the surrounding inflammatory milieu. Consistent with this hypothesis, CD4⁺ T cells do not have to be viral antigen specific to cause HSK, although virus-specific CD4⁺ T cells may initiate the process (110–114). While a variety of other mechanisms have been suggested, including auto-antigen unmasking (115) and viral molecular mimicry (116), these hypotheses fell out of favor after it was shown that the peptides proposed to produce autoreactive CD4⁺ T cells do not induce HSK in mice and are not recognized by T cells isolated from patients with HSK (114, 117).

Results of adoptive-transfer experiments performed in our laboratory with WT and HVEM KO mice indicate HVEM-mediated pathogenesis occurs when HVEM is present on radiation-resistant cells (44). Because they turn over with radiation and reconstitution, CD4⁺ T cells are not likely to be the HVEM⁺ cells responsible for pathology, at least in our adoptive-transfer model. However, interactions between HVEM (on other cell types) and HVEM ligands such as LIGHT, BTLA, or CD160 on CD4⁺ T cells could contribute to the development of HSK.

BTLA provides a corepressive signal on CD4⁺ T cells in *cis* or in *trans* (60, 66), but the majority of HVEM on naive T cells occurs in *cis* complexes with BTLA, with or without LIGHT as a part of the complex (118). In the *cis* complex, BTLA holds HVEM in an inactive state, preventing NF- κ B activation (118). Absence of HVEM from CD4⁺ T cells may make BTLA and LIGHT available to interact with HVEM on other cell types, such as long-lived, resident macrophages, activating NF- κ B signaling and inflammatory activity during HSK. This would be consistent with our finding that chimeras lacking HVEM from radiation-sensitive cells developed more severe disease than HVEM KO controls (44). BTLA levels have been reported to increase in the corneas of BALB/c HSV-1-infected

mice, although the identity of BTLA⁺ corneal cells and the effect that endogenous BTLA expression has on pathogenesis were not thoroughly investigated (119). In that study, intravenous treatment with a recombinant BTLA-expressing plasmid prior to and during ocular HSV-1 infection reduced HSK symptoms, the overall number of corneal CD4⁺ T cells, and the proportion of the cells expressing IFN- γ , although no experiments were performed to determine which cell types absorbed the plasmid and how these effects were mediated (119).

Levels of a specialized set of anti-inflammatory CD4⁺ T cells, FoxP3⁺ regulatory T cells (Tregs), reportedly expand during murine HSV-1 infection through gD-HVEM interactions, leading to slightly diminished corneal pathology in HVEM KO corneas late after infection (120). While this instance of decreased HSK symptoms in HVEM KO corneas conflicts with data from a multitude of studies from our laboratory (42–44), differences in viral strain and inoculation dose could produce variability in findings. Consistent with this report, our laboratory found that at 14 dpi, a small but real population of the HVEM⁺ population was CD4⁺, although we did not assess FoxP3⁺ expression in these cells. Due to the complexity of HVEM signaling and to its widespread expression on nearly every leukocytic population implicated in the development of HSK, it would be surprising if only one HVEM-associated cell type or function influenced ocular herpes immunopathogenesis. While we are confident that HVEM-mediated inflammation is the more potent effect, it is plausible that HVEM could have contradictory roles and that HVEM on Tregs may provide some relief during HSK. Further investigation of this discrepancy is required.

While less important for HSK development, CD8⁺ T cells control viral spread into the nervous system (121, 122) and suppress viral reactivation from the TG (123). The latency state, characterized by suppression of all viral products except the long noncoding RNA latency-associated transcript (LAT), occurs in the TG in humans and mice (124). LAT is not required for latency (125, 126), although LAT(–) viruses establish latency and reactivate less efficiently (127). Most human cases of HSK result from reactivation of a latent infection rather than from primary infection (128). Unfortunately, significant gaps in our understanding of latency and reactivation persist because the vast majority of studies are performed in murine models of primary infection (129). However, it has been established that CD8⁺ memory T cells infiltrate latently infected TGs in humans and mice, residing in close association with neurons (130, 131). The close association of CD8⁺ T cells, which typically express a variety of HVEM ligands (69), and neurons of the TG, which express HVEM (132), raises the question of direct effects of HVEM signaling on HSV latency. HVEM KO mice have lower rates of latency and reactivation than WT mice, although the HVEM KO strain also has lower titers in the eye initially, likely leading to decreased seeding of the TG (43, 133). Recently, investigators reported that LAT upregulates HVEM expression *in vivo* and *in vitro*, potentially through binding of the HVEM promoter by small noncoding RNAs derived from LAT (133). Increased HVEM expression could alter immune responses to reactivation; alternatively, HVEM-mediated NF- κ B activation could enhance survival of neurons undergoing reactivation. Because human disease is mostly caused by reactivation, our laboratory is actively pursuing murine models of recurrent disease (134) to study what contribution HVEM may make to this process.

HVEM SIGNALING IMPACTS A VARIETY OF HUMAN DISEASES, OFFERING TARGETS FOR THERAPY

We propose that, independently of viral entry, HVEM orchestrates an inflammatory response to HSV in the murine cornea that contributes to the extensive immune-mediated damage observed in HSK (Fig. 2). This finding has yet to be validated in human patients, a major limitation of this work. However, HVEM is implicated in a wide range of autoimmune, inflammatory, and infectious processes that impact an astonishing diversity of human syndromes (24). Because of this, the HVEM signaling network is a rich area of research for the discovery of new therapies.

Levels of soluble HVEM are elevated in the sera of patients with allergic asthma,

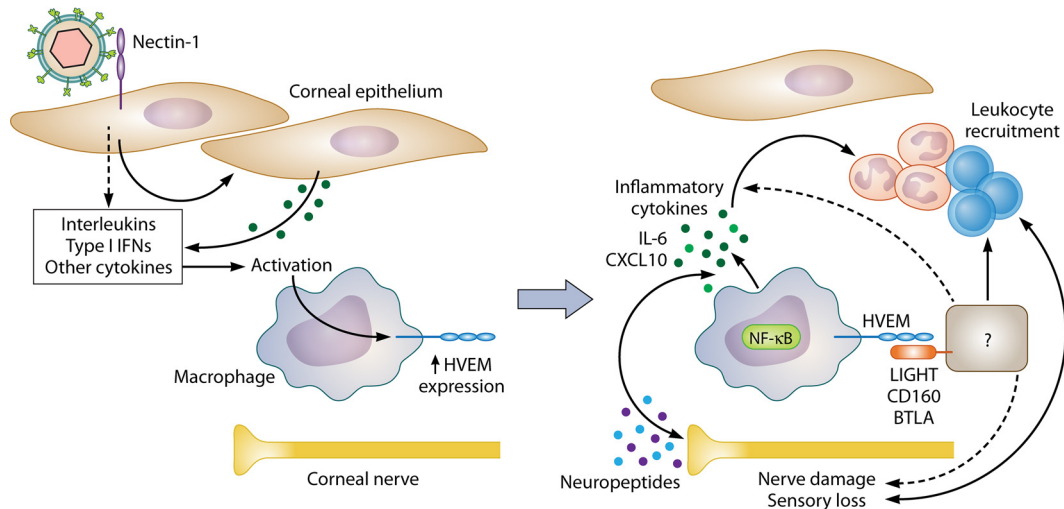


FIG 2 Proposed mechanism of HVEM pathogenesis during ocular HSV-1 infection. (Left panel) Early after infection, the majority of corneal HVEM⁺ cells is localized to CD11b⁺ Ly6C⁺ monocytic lineages rather than epithelial or endothelial cells. We propose that these HVEM⁺ cells are corneal resident macrophages, as results of adoptive-transfer experiments indicate that radiation-resistant cells are sufficient for HVEM-mediated pathogenesis and that macrophages in the cornea turn over incompletely after irradiation. HSV-1 infects the corneal epithelium, causing secretion of type I interferons and other factors that activate corneal resident macrophages, which could induce HVEM expression on these cells early after infection. (Right panel) Because the gD-HVEM interaction is dispensable for pathogenesis, we hypothesize that HVEM on corneal macrophages interacts with a host ligand on an as-yet-unidentified cell type. This results in HVEM-dependent induction of inflammatory cytokines, loss of corneal sensitivity, and immune cell recruitment. It is not yet clear whether this stems from changes in the HVEM⁺ cells operating directly, such as through NF-κB activation, or through signaling on the HVEM ligand-bearing cell. Activated macrophages may damage neighboring corneal nerves, worsening inflammation through desiccation of the cornea, or nerves themselves may influence leukocytic recruitment through release of neuropeptides. The two processes likely influence each other, synergizing to create a cycle of chronic inflammation and corneal nerve damage.

rheumatoid arthritis (RA), and atopic dermatitis (135), but HVEM/LIGHT/BTLA signaling and therapeutic opportunities within that network have been most extensively studied in models of graft-versus-host disease (GVHD) (64, 74, 136–139). GVHD is an immunologic syndrome affecting transplant recipients in which engrafted donor T cells attack host tissues, causing rampant damage (140). Investigators have had success in decreasing symptoms of GVHD in mice by targeting both BTLA-HVEM and LIGHT-HVEM interactions with blocking antibodies (137–139). Blockade of LIGHT-HVEM signaling has also been shown to increase the rate of survival of solid allografts, including pancreatic islets and cardiac transplants (141, 142). Recently, lymphoma B cells from patients with mutations in the HVEM gene (*TNFRSF14*) were found to have increased alloantigen-presenting capacity in comparison to controls, corresponding to higher levels of GVHD in patients undergoing allogeneic hematopoietic stem cell transplantation (143).

In murine experimental autoimmune uveitis (EAU), a model of human autoimmune conditions with ocular manifestations such as Behçet disease and sarcoidosis, HVEM was shown to increase disease severity by inducing pathogenic Th1- and Th17-type T cell responses (144). Similarly to HVEM KOs, LIGHT and BTLA KOs are protected from severe disease during EAU, suggesting that these ligands, in combination with HVEM, promote pathogenesis in EAU (144). If LIGHT and/or BTLA are the ligands involved in HSK as well, targeting HVEM signaling with antibodies or small-molecule inhibitors could produce novel therapies applicable to a variety of ocular inflammatory conditions.

Our laboratory has recently discovered that pathological inflammation during murine HSK can be prevented with treatment with immune-modifying nanoparticles (IMPs). Inflammatory monocytes and other circulating inflammatory cells take up IMPs and subsequently undergo apoptosis in the spleen, with the effect of diminished tissue damage under a variety of inflammatory conditions (145, 146). In our primary murine HSK model, IMP treatment improved survival of BALB/c mice, which are highly suscep-

tible to CNS involvement and mortality. It also improved corneal blink responses and decreased the levels of a variety of myeloid and lymphoid populations in the corneas of C57BL/6 mice during the chronic inflammatory phase (55). Although IMP treatment does not target HVEM signaling specifically, it achieves many of the same outcomes observed in the HVEM KO, likely by functioning downstream of HVEM, preventing HVEM-mediated recruitment of leukocytes to the cornea and the subsequent damage they cause.

CONCLUSIONS AND FUTURE DIRECTIONS

Based on previous findings and our most recent data, we propose that the presence of HVEM on corneal resident macrophages is critical for HSK development, operating either by interacting with HVEM ligands on other cells or through increased cytokine production, perhaps via NF- κ B activation (Fig. 2). We hypothesize that the presence of HVEM on these cells leads to increased corneal nerve damage, immune cell recruitment, and overall severity of disease. Because resident macrophages and nerves are physically associated in the peripheral stroma (147), it is possible that HVEM-mediated secretion of damaging cytokines from macrophages hastens corneal nerve damage, decreasing blinking and desiccating the cornea (148). The presence of HVEM on other immune cell types, including CD4⁺ T cells and PMN, is not likely to be necessary for HSK development, in light of data from adoptive-transfer experiments, but may still be contributory. Resolving pathogenic versus protective functions of HVEM, i.e., on corneal resident macrophages compared to Tregs, is an important next step for the field, as this information is critical for targeted therapy design.

There is some evidence that HVEM contributes to latent infections as well (43, 133). Whether this is physiologically significant and how this effect comes about are intriguing lines of investigation for the future. Murine models of HSK are generally based on primary infection, because, unlike humans, mice do not undergo efficient spontaneous reactivations (129). The absence of data demonstrating how the immune responses that occur during HSK differ between a primary infection and a reactivation from latency represents a significant gap in our knowledge. Other areas in need of further study are the expression and role of the host HVEM ligands during HSV-mediated pathogenesis. Studies performed with the HVEM-entry-null Δ 7-15 mutant indicated that the gD-HVEM interaction is not required for the inflammation HVEM causes; therefore, attention must be turned to the HVEM ligands BTLA, CD160, LIGHT, and LT α . To our knowledge, only BTLA expression has ever been examined in the cornea (119); expression of these molecules in the cornea is under active investigation by our laboratory. Beyond informing our mechanistic understanding of HVEM signaling during ocular herpetic infection, this information will also facilitate the testing of antibody or small-molecule therapies targeted to pathological HVEM signaling. Finally, the vast majority of studies reviewed here were performed in murine models, and studies with human tissue are needed to corroborate and expand these findings.

Both HSK signaling and HVEM signaling are complex processes involving nearly every type of leukocyte; consequently, untangling the role the HVEM plays during development of HSK has been and will continue to be a challenge. Advances made in investigation of the molecular signaling mechanisms of HVEM and its ligands will be extremely useful going forward, but the complications of bidirectional signaling interactions, with complementary or contradictory messages being delivered at the same time, remain. HVEM signaling warrants further investigation specifically in the context of immunomodulation in the eye, as our research clearly demonstrates a pathogenic and inflammatory role during ocular HSV infection. With luck, these discoveries will be translatable to new therapies for patients with this blinding condition.

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