MINIREVIEW

Virus Cell-to-Cell Transmission[▽]

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Viral infections spread based on the ability of viruses to overcome multiple barriers and move from cell to cell, tissue to tissue, and person to person and even across species. While there are fundamental differences between these types of transmissions, it has emerged that the ability of viruses to utilize and manipulate cell-cell contact contributes to the success of viral infections. Central to the excitement in the field of virus cell-to-cell transmission is the idea that cell-to-cell spread is more than the sum of the processes of virus release and entry. This implies that virus release and entry are efficiently coordinated to sites of cell-cell contact, resulting in a process that is distinct from its individual components. In this review, we will present support for this model, illustrate the ability of viruses to utilize and manipulate cell adhesion molecules, and discuss the mechanism and driving forces of directional spreading. An understanding of viral cell-to-cell spreading will enhance our ability to intervene in the efficient spreading of viral infections.

Viruses are small protein capsids that harbor genetic information. In the case of enveloped viruses, an additional lipid bilayer surrounds the capsid. In order to replicate, viruses are completely dependent on their host. They replicate their genetic information within cells, assemble and release viral progenies to infect additional cells, and spread the viral infection. The viruses discussed in this review are enveloped. They pinch off from the producer cell and enter the next cell by membrane fusion. Enveloped viruses can spread via two distinct routes, either through the cell-free aqueous environment or, alternatively, by remaining cell associated and being passed on by direct cell-cell contact. The latter mode of spread is often designated cell-to-cell transmission (for an excellent review, see reference 108). In our review, we will concentrate on lessons learned, particularly in the retrovirus field, and include cross references to other viruses.

Over the years, strong evidence has accumulated that many viruses can efficiently spread by direct cell-cell contact (57, 95, 108). An early striking observation was that viruses were able to spread in the presence of neutralizing antibodies that completely blocked the spread of cell-free virus (14, 39, 47, 82, 95). Herpes-, rhabdo-, and measles viruses were noted to spread along neuronal networks, which implied that these viruses can be transmitted via neurological synapses (54, 67). The ability of vaccinia and African swine fever viruses to induce actin tails in infected cells suggested a spreading mechanism that is related to that of the bacteria *Listeria* and *Shigella* (23, 36, 62). For many other viruses, the evidence that cell-cell contact plays a role was initially more indirect. The observed poor infectivity of cell-free virus often could not explain its rapid spreading in culture (9, 29). Early electron micrographs of HIV accumulat-

ing to high numbers at the interface between cells had a lasting impact (95). Finally, few could deny the persuasive power of time-lapse movies that directly displayed the movement of viruses from one cell to another (51, 56, 112).

Another influential observation in this field was the finding that the addition of a few dendritic cells (DC) to a culture of T cells could dramatically enhance HIV infection of T cells (18). DCs were later shown to capture and present HIV to T cells in a process that was reminiscent of how antigen-presenting cells (APC) present antigen (40, 79). Thus, following the early realization that neurotropic viruses spread along neurological synapses, a model in which immunotropic viruses such as HIV may utilize immunological synapses for efficient cell-to-cell spread began to emerge (79).

The next fundamental step in the understanding of virus cell-to-cell transmission came with the realization that HIV- and human T-lymphotropic virus (HTLV)-infected cells could establish similar cell-cell contacts between infected and uninfected T cells (53, 58, 60). T cells normally do not form long-lived immunological synapses with each other. Consequently, in analogy to immunological synapses, these cell-cell contacts were designated infectious, virological, or viral synapses (4, 53, 79). Similar observations were made for murine leukemia virus (MLV) in cultures of fibroblasts (56, 112). MLV infection established cell-cell adhesions with uninfected neighboring cells due to interactions between the viral envelope glycoprotein expressed in the infected cell and the viral receptor expressed in the target cell (112). Thus, it appears that viruses can either utilize existing cell-cell contacts or exploit basic cell adhesion biology to deliberately establish contact between infected and uninfected target cells for the purpose of efficient spreading.

VIRAL UTILIZATION OF EXISTING CELL-CELL CONTACTS

Cell-cell adhesion and specialized biological synapses are essential building blocks in tissues and organs of multicellular

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organisms (133). In order to overcome these barriers, many viruses have evolved into perfect insiders of cell-cell adhesion and biological synapses. The ability of neurotropic viruses to spread via neurological synapses was recognized many years ago and has been elegantly exploited to map neuronal circuits (27, 71, 104). Herpesviruses can spread in both directions along neurons, and their capsids can undergo anterograde and retrograde transport along microtubules (114). Spread via synapses requires that the chosen receptors specifically localize to the synapse. In fact, receptor choice allows first insights into the mechanism of virus spreading. Nectins, used as receptors by a number of herpesviruses, may localize to puncta adherens that provide a peripheral structural framework for neurological synapses (122). The discovery of tight junction components claudin-1 and occludin as hepatitis C virus (HCV) entry factors points to the use of tight junctions during HCV spread within the polarized liver epithelium (33, 98). Spread via tight junctions is common for viruses that infect epithelial layers (8, 11). Recently developed *in vitro* culture systems will allow experimental access to studying the mechanism by which these viruses spread from cell to cell (20, 61, 106).

Retroviruses and other immunotropic viruses often utilize immunological synapses for cell-to-cell spread. Contact between antigen-presenting cells and T cells is initiated by an interaction between intercellular adhesion molecules (ICAMs) expressed on APCs and LFA-1, the CD11α/CD18 αL/β2 integrin. Following the initial contact, major histocompatibility complex (MHC) molecules carrying antigenic peptides and the T-cell receptor move into the center of the contact zone, while the integrin forms an outer ring (44, 52). Talin binding to the cytoplasmic tail of integrins marks the outer rings. A center-directed flow driven by actomyosin lies behind these reorganizations (63, 90, 131). The central and peripheral zones are called the central and peripheral supramolecular activation clusters, cSMAC and pSMAC, respectively (84) (Fig. 1A). In addition to the actin-driven surface movement, cell-cell adhesion positions the PAR complex at the cytoplasmic face of the cell-cell contact zone to establish cellular polarity (72, 123). The microtubule organizing center (MTOC) moves toward the contact zone, and exocyst positioning polarizes vesicle release toward the contact zone (25, 65, 66, 75, 99, 119, 120). While these events have been described in great detail for the T-cell side, polarization is often observed on both sides of the synapse (10, 15). The duration, stability, and turnover of immunological synapses are under the control of many regulatory and inhibitory adhesion proteins (26, 123). Chemokine receptors, including HIV coreceptors CXCR4 and CCR5, serve as regulatory receptors and are recruited to immunological synapses (16, 21, 83). The concentration of both receptors in the cell-cell interface reduces the responsiveness of T cells to soluble chemokines, thereby preventing their migration and stabilizing the immunological synapse (83).

Amazingly, all three of these structural elements within the immunological synapse, cSMACs, exocytic sites within the central zone, and pSMACs, have been associated with the transmission of HIV (Fig. 1A). The relocation of HIV from vesicles or deep invaginations found in infected macrophages or dendritic cells to contact sites formed with neighboring lymphocytes was observed by time-lapse microscopy (12, 43, 136). Other HIV and HTLV accumulations between lymphocytes that appear either as buttons or outer rings are more consistent with the architecture of cSMACs and pSMACs, respectively (51, 53, 102). Mechanistically, the accumulation of viruses in the synapse could be the consequence of either *de novo* assembly in the cell-cell interface or actomyosin-driven surface movement of completely assembled particles from surrounding areas (55) (Fig. 1B). Moreover, a role for extracellular matrix components, such as galectin and collagens, has been described for the surface transmission of HTLV (92) (Fig. 1B).

While viruses utilize some features of cell-cell communication, they suppress others. For instance, HIV hijacks the cellular pathway for the presentation of antigen in order to spread the infection, but can prevent the actual presentation of antigens. This occurs through the expression of Nef, which downregulates MHC molecules (34). Nef alleles from various lentiviruses also downregulate CD4 and CD3 and remodel the actin cytoskeleton, which influences the duration and signaling events within the immunological synapse (5, 34, 103, 121).

Finally, while the exploitation of existing synapses allows efficient spreading from cell to cell, viruses also hijack the capacity of certain cell types to mediate long-range transport throughout an organism. The ability of neurotropic viruses to spread across neurons allows these viruses to enter peripheral neurons yet quickly reach the central nervous system (24, 105). Likewise, the infection of peripheral dendritic cells allows HIV to move with these cells into the lymph node for transfer to T cells (40, 76, 96). Thus, viral exploitation of existing forms of cell-cell communication extends beyond just the cellular transmission aspects and explains spreading within an organism.

CAMS AND VAMS: VIRAL MIMICRY OF CELL ADHESION MOLECULES

In addition to utilizing existing cell-cell contacts, viruses can deliberately establish cell-cell contact between infected cells and uninfected target cells (53, 58, 112). To understand the biology of this process, it is helpful to examine biological synaptogenesis. Despite the complexity of each biological synapse, synaptogenesis is initially driven by a few dominant cellular adhesion molecules (CAMs) (3, 133). These proteins alone, if expressed in cells that lack most other adhesion proteins, have the ability to form synapses and induce cellular polarity. In the case of neurosynaptogenesis, dominant adhesion activity has been attributed to the homotypic adhesion molecule SynCAM as well as the heterotypic adhesion molecule neuroligin/neurexin (13, 109). E-cadherin and ICAMs/LFA-1 are dominant homotypic and heterotypic cellular adhesion molecules that can drive the formation of epithelial and immunological synapses, respectively (63, 132). Synaptobiogenesis often proceeds through filopodial and dendritic intermediates (35, 64, 89, 113, 127, 137). Cytoplasmic signaling events downstream of cell adhesion allow the reorganization of the underlying cytoskeleton, e.g., by suppressing filopodial formation in order to organize a smooth and broad cell-cell interface (1, 118a). The establishment of cell-cell adhesion is often followed by an induction of cellular polarity that leads to the reorientation of the MTOC as already mentioned previously for the immunological synapse (31, 133).
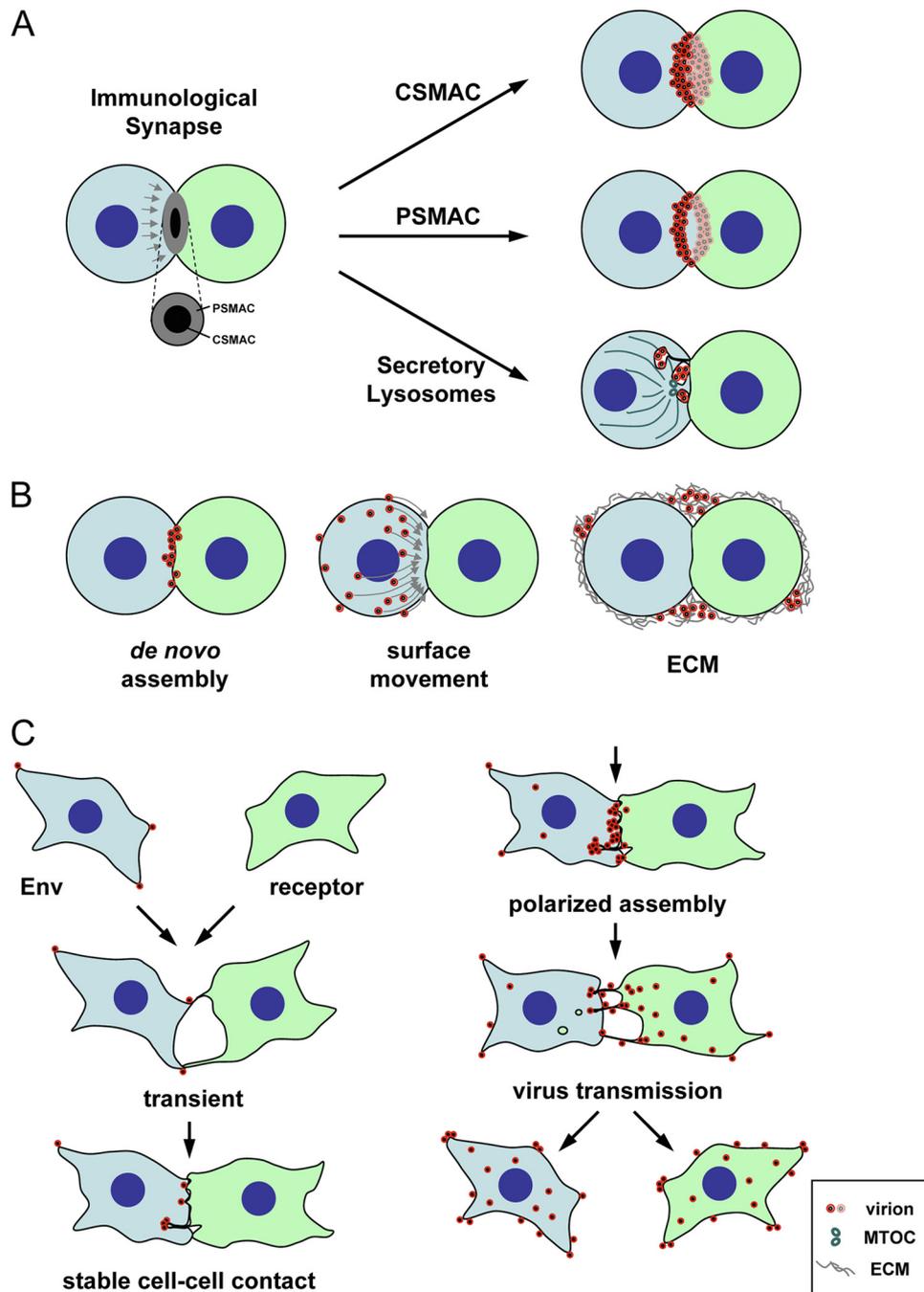
FIG. 1. Viruses can either utilize existing synaptic contacts or establish contact between infected and uninfected cells to promote viral spreading. (A) Utilization of structural elements of immunological synapses for viral spreading. The architecture of the immunological synapse consists of central and peripheral supermolecular adhesion complexes, cSMAC and pSMAC, respectively (84). The antigen-presenting cell is shown in green, and the effector cell is shown in blue. Exocytosis of secretory lysosomes can be observed in the cSMAC zone. Interestingly, all three structural elements have been associated with the accumulation and transmission of viruses from infected to uninfected cells (43, 51, 102). (B) Several distinct mechanisms can contribute to the accumulation of viruses at the synapse. The association of viruses with the cell-cell interface could be the consequence of either *de novo* assembly or surface transmission (55, 56). Extracellular matrix components (ECM) have also been observed to play a role in a peripheral mode of transmission (92). Virus-infected cells are blue, and the receptor-expressing target cells are labeled as green. (C) Establishment of adhesion between infected and uninfected cells using MLV as the model system (56, 112). MLV-infected fibroblasts were observed to establish contact with uninfected cells in a reaction driven by Env-receptor interactions. Initial transient contacts are stabilized, and virus assembly was observed to be redirected toward sites of cell-cell contact after an ∼30-min delay. Following virus transmission, cell-cell contacts are downregulated. Virus-infected cells are blue, and the receptor-expressing target cells are green.

Importantly, viruses use several mechanisms to exploit cellular adhesion biology to establish contact between the infected and uninfected target cell. First, HTLV-1 infection can upregulate endogenous adhesion proteins, such as ICAM-1, as well as components of the extracellular matrix (38, 92). Second, mouse mammary tumor virus expresses a superantigen that causes proliferation of infected B and T cells but may also stabilize and prolong contact between B cells and T cells (42, 48). Third, some viruses appear to express dominant adhesion proteins (20, 53, 58, 112). In the case of MLV, the viral protein that exhibits adhesion molecule features is the viral glycoprotein Env (112). Cells that express MLV Env can establish cell-cell contact with cells expressing the viral receptor (Fig. 1C). Env mutants that no longer interact with the viral receptor with high affinity fail to establish cell-cell contact (112). Moreover, in analogy to cellular adhesion and subsequent polarization, the establishment of adhesion between MLV-infected cells and uninfected target cells is followed by the polarized assembly of viruses at the cell-cell interface (56) (Fig. 1C). This polarized assembly process depends on signaling via the cytoplasmic tail of Env. Upon incorporation of MLV Env into budding virions, the cytoplasmic tail is cleaved off by the viral protease, thereby transforming an adhesion protein into a highly fusiogenic fusion protein that facilitates virus entry into the neighboring cells (22, 45, 101). Thus, the Env fusion protein initially functions as a viral adhesion molecule (VAM) mimicking the behavior of a cellular adhesion molecule. It will be important in the future to understand the specific adhesion biology utilized by various viruses.

While these MLV spreading experiments have been performed with transformed cancer cells that have lost many of their endogenous adhesion proteins, in lymphocytes and particularly in primary cells, VAMs likely synergize with endogenous CAMs. Indeed, the study of HIV and HTLV spreading suggests that Env from these viruses synergizes with LFA-1 and ICAM-1 but is not essential for the establishment of cell-cell contact (19, 43, 102, 118). In contrast, ICAM-1 and LFA-1 support efficient spreading, underscoring the synergy between VAMs and CAMs (49, 59, 102, 129).

Finally, virus spreading in primary cells can be distinct from that observed with tissue culture cells. In migrating primary T cells, which are already polarized prior to the establishment of cell-cell contact, HIV assembly is directed to the uropod (19) (Akira Ono, personal communication). Upon contact with uninfected cells at the leading edge, the cell turns around to establish prolonged contact through the uropod that also harbors many adhesion proteins and signaling components of the immunological synapse (6, 72, 107).

Thus, while viruses can utilize existing synapses for cell-to-cell spread, VAMs can also specifically promote adhesion between infected and uninfected cells in cultures that usually do not form synaptic contacts. To mention one more example in support of this model, while herpesviruses can spread through neurological synapses, they can also establish new cell-cell contacts outside the synapse to promote viral spread to neighboring cells (28, 108). The ability to use viral glycoprotein/receptor pairs with dominant adhesion features will also allow viruses to manipulate existing synapses (78, 126).

MECHANISM OF VIRUS CELL-TO-CELL TRANSMISSION

Viruses can spread via either a cell-free mode or a cell-associated mode involving direct cell-cell contact. Efficient virus spreading can be achieved by either route. For cell-free aqueous spreading to be efficient, a virally infected cell would have to release large numbers of viral particles and reach distant areas by diffusion. These particles must be sufficiently stable, not quickly cleared and, importantly, still able to efficiently bind and infect uninfected target cells. In contrast, if any of these criteria are not fulfilled, spreading via the cell-free mode is impaired. For instance, if viruses are not efficiently released into the extracellular milieu, spreading by the cell-free mode would obviously be blocked. However, virus retention on producer cells may not necessarily block cell-to-cell transmission. In fact, studies showing that surface retention on producer cells plays an important role in the spreading of viruses by cell-cell contact are emerging (92, 111). Intriguingly, it remains to be determined if tetherin, a recently identified antiviral factor that retains HIV particles on the cell, will promote or block cell-to-cell transmission (46, 86, 87, 94, 128). Second, released viruses may be too unstable to allow for cell-free spreading but be able to undergo rapid spreading via sites of cell-cell contact. As an alternative, released viruses could be captured and stabilized by cell surface or extracellular matrix components (85, 92, 93). Third, viral gene expression may be too low in certain cell types to allow efficient particle generation. However, inefficient virus assembly and release can be rescued at sites of cell-cell contact by locally enhancing virus assembly and release (56, 57). Fourth, cell-free viruses may not be able to efficiently bind to target cells, which likely represents one of the most commonly observed blocks to cell-free virus spreading (91, 97, 124). Single viruses, due to their small size and limited number of glycoproteins, may only weakly bind to target membranes. They may not recruit receptor to levels sufficient to induce uptake and/or conformational changes needed to infect cells. Moreover, single viruses may not be able to activate quiescent cells sufficiently to support viral replication. In contrast, Env highly enriched at broad cell-cell interfaces may recruit receptor and signaling proteins to levels that make cells susceptible to viral replication (2, 7, 125, 134, 135). It should be noted that certain extracellular components may similarly promote efficient virus binding and infection of cells (85).

These general considerations already point to a number of conditions under which viruses may not be fit enough for cell-free virus spreading. However, spread by direct cell-cell contact is likely more than a salvage pathway for the unfit. While researchers were forced to discover the importance of cell-to-cell spread for viruses with poor infectivity-to-particle ratios, it is worthwhile to consider the possibility that even the most stable viruses use cell-to-cell spread. There are several appealing advantages associated with direct cell-to-cell spread that could be exploited by many viruses. The first is speed: rather by going through all the steps of cell-free transmission, the entire extracellular replication cycle of release, transmission, and entry can proceed quickly at sites of cell-cell contact and exploit cytoskeletal forces for the purpose of spreading. Moreover, the observed enhancement of budding at sites of cell-cell contact can promote spreading at lower levels of gene expression (56).

The second is immune evasion: limited exposure time to the extracellular space can allow evasion of neutralizing antibodies (39, 51). Third, exploiting cell-cell communication is an effective way to overcome the various physical and immunological barriers within an organism in order to spread the infection.

Finally, while cell-to-cell spread has its appeal, under some circumstances, cell-free virus spread might be more advantageous. Cell-free virus is not restricted to specific cell-cell interactions and may facilitate spread from person to person. As such, it is possible to imagine that some viruses, notably HIV, may have come up with mechanisms to switch between cell-free and cell contact-dependent modes of spreading.

DRIVING FORCES FOR DIRECTIONAL VIRAL SPREAD

The spread of viral infection depends on the directional transmission of virus particles from infected cells to uninfected target cells. The most critical prerequisite for virus spreading, irrespective of whether transmission occurs via the cell-free or cell-cell contact-dependent mode, is that the viral receptor is downregulated in infected cells. Downregulation of the viral receptor in the producer cell and high affinity for the receptor expressed on the target cell alone establish an affinity gradient that drives viral spreading. This simple model is modified in an interesting way by a frequently observed retention of viruses on the surface of the producer cell despite the completion of assembly (86, 92, 111). As mentioned above, this prevents release into the medium, but it does not necessarily prevent cell-to-cell spreading (92, 111). In fact, if diffusion coefficients are comparable, diffusion along a two-dimensional surface would promote viral spread more efficiently than diffusion in three dimensions. Interestingly, following downregulation of the high-affinity receptor, viruses can be retained on the surface of their own producer cell due to nonspecific virus-cell interactions (111). Over time and due to substrate and/or extracellular matrix trapping, this retention can result in an accumulation of viruses that are subsequently efficiently transferred to target cells (Fig. 2A).

Importantly, upon contact with the target cell, viruses bind to their viral receptor with specificity and high affinity. High-affinity interactions between ligands, including viruses and their receptors at the plasma membrane, can result in the establishment of a link from the surface to the underlying actin cytoskeleton (68, 69). Filamentous actin (F-actin) undergoes constant turnover that is driven by assembly "pushing" at the tip and myosin II-driven "pulling" at the base of the actin filament (80). Consequently, viruses end up "surfing" toward endocytic hot spots at the cell body (17, 68–70, 110, 115). As such, viruses do not recruit an individual myosin motor to each particle but, rather, utilize the general turnover of F-actin. Thus, viruses engage a high-affinity interaction with receptor-expressing cells that allows them to utilize actin-driven motion to move toward target cells (17, 111, 112) (Fig. 2B).

This affinity-based mode of transmission has recently been reported for transient interactions between fibroblasts chronically infected with MLV and uninfected target cells (111). In contrast to transient cell-cell interactions, prolonged interactions between infected and uninfected cells lead to the anchoring of target cell membranes, including filopodial membranes, in the infected cell (56, 112). The infected cell in turn redirects
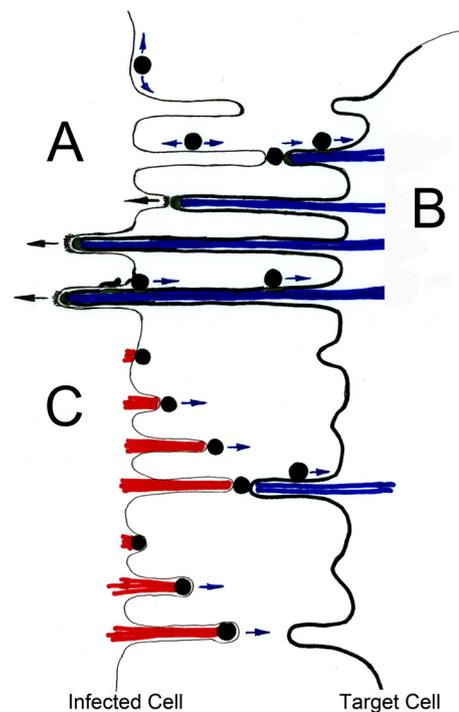

FIG. 2. Driving forces for directional spreading of viruses (adapted from reference 113 with permission of the publisher). (A) Affinity-based model for directional spreading. Downregulation of the viral receptor in the infected cells allows the establishment of a general affinity gradient between the infected and uninfected cell. Additional virus interactions with the infected cell can retain completely budded viral particles at the surface of the infected cell. Low-affinity retention on the surface of producer cells allows for diffusive movement along the cell surface (111). This process can position viruses for subsequent high-affinity interactions with target cells expressing receptor, thereby driving directional spreading of surface associated viruses. (B) Viral transmission driven by retrograde F-actin flow of the target cell. High-affinity interactions with receptor (green) in the target cell allows surface-associated viruses to engage the underlying F-actin flow (blue) to move toward the cell body of the target cell (17, 68, 69, 110, 115). This process, originally designated "virus surfing," can promote a driving force for cell-to-cell spreading for the surface-retained viruses presented in panel A. In addition, long-term Env-receptor interactions between infected and uninfected cells can anchor target cell membranes directly at the infected cells (112). Because virus assembly can be redirected toward these sites of cell-cell contact (56), viruses can immediately engage target cell F-actin flow (blue) to move toward the target cell upon completion of assembly. (C) Viral transmission driven by actin assembly in the infected cell. In contrast to the depictions in panel B, viruses can induce actin comet tail formation (red) inside the infected cell to propel themselves toward neighboring cells (23, 30, 62). While vaccinia virus induces actin tails (red) beneath the plasma membrane of completely released viruses, capsids of the African swine fever virus induce actin tails (red) while still topologically within the cytoplasm. Such actin-propelled motion may be sufficient to drive transmission, but a combination of actin-driven movements both away from the infected cell (red) as well as toward the target cell (blue) are also plausible (81).

virus assembly directly to these sites of cell-cell contact (56). Consequently, as soon as viruses assemble, they can utilize retrograde F-actin flow to propel themselves toward target cells. Thus, both transmission pathways exploit actin dynamics from target cells for efficient cell-to-cell transmission (Fig. 2A and B).

In contrast to this process, other viruses induce actin tails within the infected cell to propel themselves toward target cells (23, 30, 62). Processes with two distinct membrane topologies have been observed. In the case of vaccinia virus, the virus is already completely released, but it uses the proteins A33 and A36 to induce the assembly of actin tails that propel viruses toward neighboring cells (17, 23, 37) (Fig. 2). In contrast to the case with vaccinia virus, African swine fever virus capsids form actin tails when they are still within the cytoplasm (62) (Fig. 2). Thus, viruses have evolved clever ways to utilize the actin cytoskeleton in either infected cells or target cells to propel themselves toward neighboring cells.

In addition to these fundamentally distinct mechanisms of cell-to-cell spread, viral spreading via the interior of nanotubes has been proposed but requires validation (32, 41, 113). Given that the transfer of capsid remains Env and receptor dependent, viruses such as HIV may move along the outside of nanotubes, not unlike MLV along filopodial bridges (118).

MIND THE GAP: THE EXPERIMENTAL APPROACH TO CELL-TO-CELL TRANSMISSION

Having discussed the mechanism of cell-to-cell transmission, we turn to the question of how a researcher can experimentally address whether a cell-free or cell-associated mode of virus spread is used by a given virus. This is more complicated than previously thought, because both modes usually involve extracellular virus. Most of the existing assays are not very convincing, and there is a need to combine functional and imaging approaches to collectively build an argument. The first experimental hint that cell-to-cell transmission may play a role is often that the infectivity produced by infected cells and released into the culture supernatant does not match the observed infectivity when the same number of infected cells is cocultured with uninfected cells (9). Second, fast kinetics of spreading in cocultures compared to that of cell-free virus has provided an argument for direct cell-to-cell spread (9, 29, 30). Third, viral growth in plaques, while potentially indicating direct cell-to-cell spread, may just reflect the decline of diffusion that is proportional to the square of the distance. Thus, several experimental conditions to suppress either diffusion or to prevent cell-cell contact have been explored. Diffusion has been slowed using viscous materials, such as methylcellulose (56, 130). Direct cell-cell contact has been prevented by coculturing infected and uninfected cells in Transwells, porous membranes that while preventing cell migration, allow virus diffusion (51, 74). Alternatively, shaking of cell cultures is used to prevent the formation of stable cell-cell contacts (74, 117).

Some of the most convincing experimental evidence in support of cell-to-cell transmission is resistance to neutralizing antibodies that completely block cell-free virus (14, 39, 47, 82, 95). These examples are most frequently observed for cell-cell contacts of a broad nature that are known to be tight (39). In contrast, many cell-cell contacts are less tight and remain sensitive to neutralizing antibodies (74, 112).

In the light of these experimental difficulties, morphological and imaging approaches have been very helpful. The accumulation of viruses, potentially even budding sites, specifically at sites of cell-cell contacts, can provide an intriguing snapshot that suggests the existence of a dynamic process (9, 58, 79, 95).

The "seeing is believing" approach was most convincingly demonstrated in time-lapse microscopy videos that directly monitored the movement of viruses from one cell to another at sites of cell-cell contact (51, 56, 112). Moreover, the transfer of particles apparently correlated with the subsequent infection of these cells (51). Single-cell imaging that incorporates functional readouts into the imaging approach can generate convincing data. The correlation of live-cell imaging with scanning electron microscopy and/or tomography as well as super-resolution fluorescence can provide detailed mechanistic insights, such as whether viruses move along the surface or through the interior of cellular bridges (50, 73, 74, 92, 102, 112).

One of the critical challenges for the field of cell-to-cell transmission is to understand how viruses actually spread *in vivo* in living organisms. In addition, it will be critically important to identify cellular factors and small-molecule inhibitors that specifically affect one or the other mode of transmission. Early successes are the specific dependence of cell-cell contact-mediated spreading, but not that of cell-free virus on contributions from the cytoskeleton (51, 53, 58, 77). The identification of Zap70 and abl tyrosine kinases represent the first cellular factors required specifically for the contact-mediated spread of HIV and vaccinia virus (88, 100, 116). Hopefully over the next years, the field will identify several molecules specifically required for the contact-mediated transmission of viruses. Novel reporter constructs that specifically measure only contact-mediated spread will greatly facilitate this approach for retroviruses (77). The identification of such factors would also provide novel targets for antiviral therapies designed to block virus cell-to-cell transmission.

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