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Pedro M. Matos, Emory University
Mariana Marin, Emory University
Byungwook Ahn, Emory University
Wilbur Lam, Emory University
Nuno C. Santos, Universidade de Lisboa
Gregory Melikian, Emory University

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Anionic Lipids Are Required for Vesicular Stomatitis Virus G Protein-mediated Single Particle Fusion with Supported Lipid Bilayers*§

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Pedro M. Matos§¶1, Mariana Marin1, Byungwook Ahn1, Wilbur Lam1, Nuno C. Santos1, and Gregory B. Melikyan*§

From the †Emory Children’s Center, Department of Pediatrics, Emory University School of Medicine, Atlanta, Georgia 30322, the ‡Instituto de Medicina Molecular, Faculdade de Medicina, Universidade de Lisboa, Av. Prof. Egas Moniz, 1649-028 Lisbon, Portugal, and the §Children’s Healthcare of Atlanta, Atlanta, Georgia 30322

Background: Regulation of virus entry by host lipids is poorly understood.

Results: Imaging of low pH-mediated fusion between single vesicular stomatitis pseudoviruses and lipid bilayers revealed a striking reliance on anionic lipids.

Conclusion: The dependence of fusion on late endosome-resident anionic lipids suggests a new means for regulating the virus entry sites.

Significance: Reliance on specific lipids for fusion may shed light on future antiviral strategies.

Viral glycoproteins mediate fusion between viral and cellular membranes upon binding to cognate receptors and/or experiencing low pH. Although activation of viral glycoproteins is thought to be necessary and sufficient for fusion, accumulating evidence suggests that additional cellular factors, including lipids, can modulate the fusion process. Understanding the role of lipids in virus entry via endocytosis is impeded by poor accessibility and the highly diverse nature of endosomes. Here we imaged fusion of single retroviral particles pseudotyped with the vesicular stomatitis virus (VSV) G protein with dextran-supported lipid bilayers. Incorporation of diffusible fluorescent labels into the viral membrane and the viral interior enabled detection of the lipid mixing (hemifusion) and content transfer (full fusion) steps of VSV G-mediated fusion at low pH. Although single virus fusion with supported bilayers made of zwitterionic lipids could not be detected, inclusion of anionic lipids, phosphatidylserine, and bis(monoacylglycerol)phosphate (BMP), greatly enhanced the efficiency of hemifusion and permitted full fusion. Importantly, lipid mixing always preceded the opening of a fusion pore, demonstrating that VSV G-mediated fusion proceeds through a long-lived hemifusion intermediate. Kinetic analysis of lipid and content transfer showed that the lags between lipid and content mixing defining the lifetime of a hemifusion intermediate were significantly shorter for BMP-containing compared with PS-containing bilayers. The strong fusion-enhancing effect of BMP, a late endosome-resident lipid, is consistent with the model that VSV initiates fusion in early endosomes but releases its core into the cytosol after reaching late endosomal compartments.

Entry of enveloped viruses into a host cell is a complex, multistep process mediated by viral glycoproteins, which are activated upon encountering cellular receptor(s) and/or low pH (1,2). Generally, the entry of enveloped viruses proceeds through attachment to a host cell, binding to cognate receptor(s), and either fusion of the viral membrane with the cell plasma membrane or virus internalization and low pH-dependent fusion with endosomes (3). Recent evidence suggests that viruses that do not rely on low pH can also enter through endocytic pathways (4–7). Viral envelope glycoproteins are thought to mediate energetically unfavorable membrane fusion events by releasing the energy stored in their native structure upon refolding into a thermodynamically favorable conformation (8–11). Despite the highly diverse structures of viral fusion glycoprotein, they are categorized into three classes on the basis of their structural and biochemical properties (2,12).

VSVG protein is a class III viral fusion protein that is well characterized both structurally and functionally (13–15). Interestingly, pH-induced conformational changes in VSV G are reversible, as the protein assumes the native conformation upon returning to neutral pH in the absence of a target membrane (16–18). This unusual feature of VSV G indicates that the difference between free energies of the initial “metastable” and final “stable” conformations is not large and that, therefore, little energy can be released by this protein to drive membrane fusion (19). These considerations warrant further studies of VSV G-mediated fusion to gain critical insights into the universal principles by which viral proteins promote membrane merger.

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1To whom correspondence should be addressed: Department of Pediatrics Infectious Diseases, Emory University, 2015 Uppergate Dr., Atlanta, GA 30322. Tel.: 404-727-4652; Fax: 404-727-9223; E-mail: gmeliki@emory.edu.

2The abbreviations used are: VSV, vesicular stomatitis virus; PS, phosphatidylserine; BMP, bis(monooleoylglycerol)phosphate; SLB, supported lipid bilayer(s); VSVpp, vesicular stomatitis virus G protein-pseudotyped particle(s); Did, 1,1-di-octadecyl-3,3,3,3-tetramethylindocarbocyanine; POPE, 1-palmitoyl-2-oleoyl-sn-glycerol-3-phosphoethanolamine-N-(carboxyfluorescein); Chol, cholesterol; POPE, polyethylene glycol; mKO, CoralHue® monomeric Kusabira Orange fluorescent protein; FRAP, fluorescence recovery after photobleaching; ROI, region of interest; NC, nucleocapsid; MLV, murine leukemia virus.
Lipid Dependence of VSV G-mediated Single Virus Fusion

Despite extensive efforts, the host cell receptor(s) for VSV entry has not been unambiguously identified. Phosphatidylserine (PS) has been shown to promote VSV fusion and entry into cells (20–23). However, this lipid is unlikely to be the primary VSV receptor because VSV binding and infection do not correlate with the level of PS in the plasma membrane, and blocking PS with annexin V does not inhibit VSV infection (24). Moreover, the fact that VSV can merge with lipid vesicles made of phosphatidylcholine at low pH (21, 25) indicates the lack of reliance on a proteinaceous receptor or a specific lipid. On the other hand, lipid composition is known to modulate the kinetics and efficiency of viral fusion (26) and could, therefore, play a role in the regulation of G protein-mediated fusion. Recent studies highlight the importance of an endosome-specific anionic lipid, bis(monoacylglycero)phosphate (BMP), also referred to as lysobisphosphatidic acid (27), in promoting fusion of unrelated viruses entering cells via endocytosis (28–31). For instance, dengue virus fusion with liposomes and with cell membranes has been shown to depend on anionic lipids, including BMP (30), which is one of the major lipid species in late endosomes and multivesicular bodies (27, 32). BMP is thought to facilitate cellular entry of different pathogens by regulating back-fusion between intraluminal vesicles and the limiting membrane of an endosome (33, 34). Interestingly, a BMP-dependent back-fusion process has also been implicated in the VSV nucleocapsid release into the cytosol following virus fusion with intraluminal vesicles (28, 29). However, this two-step VSV entry model is not universally accepted because this virus appears to quickly fuse with early endosomes prior to entry into multivesicular bodies (35, 36).

Single virus trafficking in host cells is an increasingly popular tool to study the viral entry process (37). However, real-time detection of fusion events that culminate in the viral content release is technically challenging and, therefore, less common (4, 38–40). The main advantage of these approaches is the ability to study events at a single virus level, which allows distinguishing specific subpopulations of particles or characteristics hidden in bulk assay data.

Although single virus imaging in live cells provides important insights into the entry process, the complexity of vesicular trafficking and the heterogeneity of endosomal compartments impede mechanistic studies of viral fusion. Dissecting the viral entry processes in a controlled environment can shed light on the mechanism of fusion and on the host factors required for completion of this reaction. A powerful approach to address these questions is to reconstitute viral fusion in model systems such as liposomes and supported lipid bilayers (SLB). Recent advances in imaging of membrane fusion using supported lipid bilayers or tethered single liposomes have brought new mechanistic insights into the membrane merger mediated by cellular fusion proteins (41–43) and viral glycoproteins (44–46). Rapid pH-dependent hemifusion/fusion of single influenza and Sindbis viruses with SLB was observed by monitoring redistribution of a lipophilic dye incorporated into the viral membrane (hemifusion) (46) or by visualizing the transfer of both viral membrane and content markers (full fusion) (44, 45).

Here, we report the direct visualization of VSV G-pseudotyped particle (VSVpp) fusion with dextran-supported lipid bilayers. We took advantage of the murine leukemia virus (MLV) pseudotyping system, which allows incorporation of a releasable fluorescent protein marker into the virus (5) and of a membrane dye into the viral envelope. This labeling strategy permitted the detection of both lipid mixing (hemifusion) and content release (full fusion) steps of the VSVpp fusion process. Imaging of single VSVpp revealed that both hemifusion and fusion with supported bilayers were markedly promoted by anionic lipids, POPS, or BMP. Although BMP-containing bilayers supported faster conversion of hemifusion to full fusion compared with POPS bilayers, similar probabilities of fusion and identical effective sizes of nascent fusion pores were consistent with the lack of requirement for a specific lipid for productive VSV entry. The strong fusion-stimulating effect of BMP indicates, however, that this endosome-resident lipid may modulate the outcome of low pH-induced VSV fusion with intracellular compartments.

**EXPERIMENTAL PROCEDURES**

**Virus Production**—Fluorescently labeled pseudoviruses were produced in HEK293T/17 cells using PolyFect transfection reagent (Qiagen, Valencia, CA). Cells grown on 10-cm dishes were transfected with 2 µg MLV-Gag-Pol, 1 µg MLV-Gag-mKO, 3 µg pMLV-LTR-LacZ, and 3 µg of pMDG-VSV-G. Twenty-four hours post-transfection, cells were labeled with 10 µM 1,1'-dioctadecyl-3,3,3,3'-tetramethylindodicarbocyanine (DiD) for 4 h in a CO₂ incubator at 37 °C. Cells were washed, covered with 6 ml of fresh phenol red-free growth medium, and incubated for an additional 24 h. Virus-containing medium was collected 48 h post-transfection, passed through a 0.45 µm filter, aliquoted, and stored at −80 °C. The infectious titer was determined by a β-galactosidase assay in TZM-bl cells (47). Vectors expressing MLV-Gag-pol and MLV-LTR LacZ (48) were provided by Dr. Walther Mothes (Yale University). The pMDG-VSV-G expression vector was provided by Dr. John Young (Salk Institute). The construction of the MLV-Gag-mKO expression vector was described in Ref. 5.

**Preparation of Small Unilamellar Vesicles**—1-palmitoyl-2-oleoyl-sn-glycero-3-phosphocholine (POPC), 1-palmitoyl-2-oleoyl-sn-glycero-3-phospho-L-serine (POPS), BMP (SR isomer) and 1,2-di oleoyl-sn-glycero-3-phosphoethanolamine-N-(carboxyfluorescein) (CF-PE) were purchased from Avanti Polar Lipids (Alabaster, AL). Cholesterol (Chol) was from Sigma. Lipids were mixed in chloroform and dried to a thin film in a round-bottom glass flask under an argon stream. Traces of solvent were removed by placing the flask in a vacuum chamber for at least 3 h. The lipid film was hydrated in PBS buffer (Cellgro, Mediatech, Inc., Manassas, VA) to make a final lipid concentration of 1 mM. The lipid suspension was subjected to several freeze/thaw cycles and sonicated with a probe sonicator (VWR, Radnor, PA) on ice to avoid sample heating. The obtained lipid suspension was cleared by centrifugation at 10,000 × g for 5 min.

**Coverslip Cleaning and Functionalization with Dextran**—Glass coverslips (no. 1.5, 22 × 22 mm, Corning, NY) were thoroughly cleaned using the following steps: 1) immersion into hot 2% Hellmanex (Hellma Analytics, Müllheim, Germany) for 30 min, 2) sonications in Hellmanex for 15 min while warm, 3)
rinsing extensively with Milli-Q water, 4) immersion into Piranha solution (3:1 concentrated sulfuric acid/hydrogen peroxide 30% without stabilizer) for 30 min, 5) rinsing extensively with Milli-Q water, 6) sonication in Milli-Q water for 10 min, and 7) rinsing extensively with Milli-Q water. The cleaned coverslips were dried in an oven for 1 h at \( \sim 100^\circ \text{C} \).

Dextran functionalization was carried out as described previously (45). Briefly, cleaned coverslips were immersed in 0.2% (3-glycidoxypropyl)trimethoxysilane (Sigma) in isopropanol for 5 min under gentle agitation. After rinsing with pure isopropanol, the coverslips were cured for 1 h at \( \sim 80^\circ \text{C} \). A 30% (w/v) dextran (\( M, 5 \times 10^6 \), Sigma-Fluka) solution was prepared using Milli-Q water, and air bubbles were removed in a vacuum chamber. The coverslips were carefully covered with dextran solution (\( \sim 1 \text{ ml} \)) and incubated in a humidified chamber for \( \sim 24 \text{ h} \). Excess dextran was rinsed off by dipping repeatedly in water and leaving the coverslips in water for 2 days. The coverslips were dried in an oven for 1 h at 80 \(^\circ\text{C}\) and stored in a vacuum desiccator.

**Microfluidic Chamber Design and Fluorescence Microscopy**—The microfluidic polydimethylsiloxane (PDMS) chamber was fabricated using both standard photolithography and soft lithography. To prepare the soft mold for the microfluidic channels, a negative photoresist (MicoChem, SU-8) was spin-coated and patterned on a 4-inch silicon wafer. The patterned wafers were silanized using hexamethyldisilazane so that the wafers were dried in an oven for 1 h at 80 \(^\circ\text{C}\) and stored in a vacuum desiccator.

The microfluidic PDMS chamber had two 10-mm grooves that were 1 mm wide and 75 \( \mu \text{m} \) deep. Small unilamellar vesicles over the hydrated dextran layer of the glass coverslip were usually formed, as observed by the fluorescence of CF-PE incorporated in the lipid mixtures. Next, SLB were tested for fluidity using FRAP (51). Briefly, a circular region of interest (radius 8 \( \mu \text{m} \)) was photobleached using a brief pulse of the 488-nm line of an argon laser set at 100% power. The fluorescence recovery was monitored by low-intensity laser excitation for at least 5 min. The fluorescence of a distant circular region of interest (ROI) served as a reference signal.

Intensity profiles for the bleached and reference ROIs were analyzed using ImageJ software. The fluorescence recovery trace was double-normalized to take into account a possible inadvertent bleaching effect upon image acquisition

\[
F_{\text{norm}} = \frac{F_{\text{Ref (pre)}}}{F_{\text{Ref (t)}}} \times \frac{F_{\text{FRAP (t)}}}{F_{\text{FRAP (pre)}}}
\]

where \( F_{\text{Ref}} \) is the mean fluorescence intensity of the reference ROI, \( F_{\text{FRAP}} \) is the intensity of the bleached ROI, \( t \) is the time after photobleaching, and \( \text{pre} \) stands for the mean intensity before photobleaching.

The normalized traces of the bleached ROI were fitted using a diffusion model for a circular spot (52)

\[
F_{\text{FRAP (norm) (t)}} = a_0 + a_1 \cdot e^{-\frac{\tau}{2(t-t_{\text{bleach}})}} \cdot \left[ I_0 \left( \frac{\tau}{2(t-t_{\text{bleach}})} \right) + I_1 \left( \frac{\tau}{2(t-t_{\text{bleach}})} \right) \right] \cdot \tau = \frac{w^2}{D}
\]

where \( t_{\text{bleach}} \) is the time of bleaching, \( \tau \) is the characteristic diffusion time, \( w \) is the radius of the ROI, \( D \) is the diffusion coefficient, and \( I_0(x) \) and \( I_1(x) \) are modified Bessel functions. Nonlinear fit of the data was performed in GraphPad Prism version 5.

The results of the FRAP assay for each type of lipid mixture used are shown in Fig. 1D, and the corresponding diffusion coefficients are in the figure legend. Diffusion coefficients around 1 \( \mu \text{m}^2/\text{s} \), corresponding to free lipid diffusion in bilayers, were deemed acceptable for virus fusion experiments. Systematic verification of the bilayer fluidity is important because a fraction of bilayers did not exhibit fluorescence recovery after photobleaching (see also Ref. 49).

**Imaging of VSVpp Fusion**—All experiments were performed at room temperature. For each imaging experiment, a sample of 5 \( \mu \text{l} \) of the virus stock solution was diluted in 100 \( \mu \text{l} \) of PBS and vortexed briefly. Approximately 90 \( \mu \text{l} \) of viral suspension was continuously flowing into a channel. Viruses adhered to the bilayer reasonably well, permitting subsequent washing with BPS for 1–2 min to remove unbound viruses. Next, image
acquisition was started, and perfusion with an MES (pH 5.5) buffer (144 mM NaCl, 2 mM CaCl₂, 30 mM MES) was initiated. This acidic buffer quenched the CF-PE fluorescence, marking the exact time of the pH change (similar to an approach introduced in Ref. 45). Images were acquired with 2-s intervals for 11 min. At the end of the acquisition, the flow channel was cleaned by flowing 1 ml of 70% ethanol at a higher flow rate before removing the chamber from the microscope.

**Image Analysis**—Hemifusion and fusion events were initially identified by visual inspection of image sequences. Next, double-labeled (DiD and mKO) viral particles were detected, and their mean fluorescence intensities over time were measured using Speckle TrackerJ, a recent freely available ImageJ plug-in (53). The program identified fluorescent particles and tracked them until the fluorescence intensity/particle size decreased below a defined threshold. DiD-only labeled particles (likely dye aggregates) were excluded from the analysis. Virus hemifusion was defined as DiD redistribution into SLBs without the loss of mKO, whereas full fusion was defined as release of DiD followed by loss of the content marker. The mean fluorescence intensity profiles for each marker (CF-PE, DiD, and mKO) were obtained using small circular regions encompassing the particles. Analysis of these profiles yielded the following key events used for kinetic analysis. The pH drop was assessed by decay of the CF-PE signal; the onset of lipid mixing was defined as the onset of DiD dequenching or, when dequenching was weak, by the onset of fluorescence decay because of DiD dilution; the onset of content release was detected by mKO fluorescence decrease; and completion of content release was defined as the time point at which the mKO signal reached the background level. The non-parametric Mann-Whitney rank sum test was used to evaluate the differences among the kinetic parameters for bilayers of different composition.

**RESULTS**

**Direct Visualization of Single VSV<sub>pp</sub> Hemifusion and Fusion Events**—The microfluidic PDMS chamber allowed for an easy solution change and for a quick formation of supported lipid bilayers, as described under “Experimental Procedures” (Fig. 1, A and B). After testing the SLB fluidity by FRAP (Fig. 1D), double-labeled VSV pseudoviruses were drawn into the chamber and allowed to attach to the bilayer for 5 min. VSV<sub>pp</sub> readily adhered to bilayers and remained relatively immobile throughout the experiments despite the solution changes. The same volume of diluted viral stock was passed through a flow channel in each experiment, resulting in adhesion of 60–100 particles/
VSVpp were colabeled with the membrane dye, DiD, and the viral content marker, the MLV-Gag polyprotein tagged with mKO (monomeric Kusabira Orange, Ref. 54), as described in Ref. 5. The Gag-mKO marker is cleaved upon MLV maturation, producing the nucleocapsid-mKO (NC-mKO) fragment, which is not associated with the viral core and is, therefore, readily released upon viral fusion (4, 5, 55). For low pH-triggered fusion, mKO is a better marker than GFP because its resistance to quenching by low pH (pK_a/Hi11349<5 (54)) minimizes the possibility that the loss of mKO signal is due to acidification of the viral interior. This strategy allowed the detection of DiD transfer into SLB as a marker for hemifusion as well as detection of the viral content release as a marker for full fusion (Fig. 1C).

VSVpp fusion with SLB was initiated by perfusion with an acidic (pH 5.5) buffer. The exact time of acidification was determined on the basis of quenching of CF-PE fluorescence, similarly to the approach described in Ref. 45. Shortly after acidification, radial diffusion of DiD into a bilayer, away from the viral particles, was observed for a fraction of the virions (Fig. 2). Typical fluorescence intensity profiles for fusion events are shown in Fig. 2, A and B, and Fig. 3. The DiD signal usually increased because of fluorescence dequenching and then decayed upon spreading into the supported bilayer. These lipid mixing events reflecting virus-SLB hemifusion were not observed at neutral pH. Following the lipid mixing events, a fraction of particles lost their content marker, NC-mKO. Because the mKO signal from neighboring particles that had no change in DiD fluorescence remained stable, the loss of the mKO signal following the hemifusion step signified the viral content release through a fusion pore (Fig. 2, D–E, and supplemental movie 1). The existence of a hydrated dextran layer supporting the membrane provided sufficient space to accommodate the viral content marker and to permit its diffusion away from a particle. The four sequential events (acidification of the flow channel, onset of lipid mixing, as well as the onset and time of completion of content release) are marked by vertical lines in Fig. 2, A and B. These four parameters were used to compare the kinetics of VSVpp fusion with SLB of a different lipid composition (see examples in Fig. 3).

Anionic Lipids Are Essential for VSV G-mediated Hemifusion and Fusion—The total number of double-labeled VSVpp attached to POPS and BMP membranes was comparable, but a
larger number of pseudoviruses adhered to POPC-based membranes lacking anionic lipids (see legend for Fig. 4A). However, despite the larger number of adhered particles, POPC bilayers did not support complete VSVpp fusion, and only 2.5% of double-labeled particles exhibited lipid mixing (hemifusion) with these bilayers (Fig. 4A). In sharp contrast, 31.1% and 41.5% of VSV pseudoviruses released DiD into POPS and BMP bilayers, respectively (Fig. 4A). BMP more potently facilitated lipid mixing than an equal amount of POPS. Under our experimental conditions, double-labeled particles released their lipid markers much more frequently than their contents. Complete fusion was detected for 6.6% of particles following their hemifusion with POPS bilayers and for 5.1% particles adhered to BMP bilayers. In other words, a significant fraction of viruses underwent hemifusion but did not form a fusion pore large enough to allow NC-mKO escape into the dextran layer underneath SLB (Figs. 1C and 2, C and F, and supplemental movie 2). Hemifusion always preceded full fusion. The fluorescence signal from...
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FIGURE 4. Analyses of the extent and kinetics of VSV pseudovirus fusion. A, the relative frequencies of fusion outcomes (no fusion, hemifusion, and full fusion) are shown for fusion of double-labeled VSVpp with SLBs of different lipid compositions. Hemifusion is defined as dequenching/loss of the DiD signal without the content release, whereas fusion is defined as the loss of both membrane and content markers (DiD and mKO, respectively). The total number of membrane-bound particles immediately after the pH drop was 398 for POPC, 257 for POPS, and 275 for BMP bilayers using four independent image fields for each lipid composition. B, distribution of the time elapsed between the pH drop (CF-PE fluorescence quenching) and the onset of lipid mixing for double-labeled particles that underwent hemifusion or fusion. The difference in the lag times before the lipid mixing for POPC and BMP was not significant (p > 0.93), but the differences between the hemifusion kinetics for these bilayers and POPS-based membranes were significant (p < 0.006 for POPS and p < 0.023 for BMP). The hemifusion half-time (t_hf) was determined by fitting the data with equation N = (A/t[H14001] + t), where A is the total number of events and t is the time. The green and red dashed vertical lines with corresponding numbers at the x axis correspond to the t_hf values for BMP and POPS bilayers, respectively.

double-labeled particles that did not exhibit lipid or content mixing was stable over the course of the experiment, demonstrating that inadvertent photobleaching of fluorophores was negligible (supplemental Fig. 1). In conclusion, 20 mol % of POPS or BMP greatly enhanced the efficiency of VSVpp hemifusion and, importantly, enabled complete fusion compared with the basal lipid mixture lacking anionic lipids (Fig. 4A).

BMP Shortens the Lag Time between Lipid Mixing and Fusion Pore Formation—As indicated above, lipid mixing events were rare for POPC bilayers lacking anionic lipids. This is in contrast with POPS or BMP bilayers supporting a much greater number of lipid mixing events. To further assess the role of anionic lipids in VSVpp fusion, we analyzed the distribution of delay times from the pH drop to the onset of lipid mixing (Fig. 4B). After acidification, VSVpp hemifusion with POPS and BMP bilayers occurred with the half-times of 44 s and 55 s, respectively (Fig. 4B). The difference in lag times to the onset of lipid mixing with BMP and POPS bilayers was not statistically significant (p > 0.93). These results suggest that although anionic lipids promote VSVpp hemifusion, this process does not require a specific lipid. It thus appears that the enhanced VSVpp fusion with bilayers containing anionic lipids is due to an electrostatic effect.

The kinetics of VSVpp fusion were analyzed in more detail using the following parameters illustrated in Figs. 2 and 5: 1) time to the onset of lipid mixing/hemifusion, 2) time to the onset of content release, 3) lag time between lipid mixing and content release for a given particle, and 4) the time from the onset to completion of the NC-mKO release as a measure of the relative size of a fusion pore. All full fusion events were preceded by lipid mixing (Figs. 2 and 3), demonstrating that VSVpp fusion proceeded through a relatively long-lived hemifusion intermediate. This analysis revealed that the distribution of the times from acidification to content release was slightly faster for BMP bilayers than for POPS bilayers, but the difference was not statistically significant (Fig. 5, A and B). Importantly, the lag time between lipid mixing and content release was significantly shorter for BMP when compared with POPS bilayers (Fig. 5, A and B). These results indicate that BMP can accelerate the conversion of a hemifusion intermediate into a functional fusion pore. Although POPS supports the efficient formation of fusion pores, this process is slower than with BMP bilayers.

We next asked whether the effective sizes of VSV G-mediated fusion pores are lipid-dependent. By analyzing the mKO release profiles for individual events on the basis of the fluorescence signal decay (Fig. 3), we found that the time required for complete release of NC-mKO varied. In many cases the mKO signal dropped virtually instantaneously (e.g. Figs. 2A and 3, A, C, and E), but there were also particles exhibiting a slow decrease of the signal (e.g. Figs. 2B and 3D and F). Thus, VSV G formed fusion pores of different initial sizes. To assess whether lipids can alter the time between the onset and the end of the mKO signal decrease, we compared the VSVpp fusion with POPS and BMP bilayers and found that the distributions of this parameter were identical (Fig. 5C). These findings argue against a specific effect of anionic lipids on the apparent size of a fusion pore.

Another prominent feature of the VSV G-mediated fusion was the two-step release of the mKO marker (Fig. 3, A and H). The two-step release of the viral content likely reflected the pore flickering. The transient halting of content release is likely due to pore closure or shrinkage, whereas resumption of release reflects subsequent pore dilatation (56, 57). Thus, fusion pores formed by VSV G in supported bilayers exhibited dynamic features similar to those formed by other viral proteins (38, 56, 58, 59).

DISCUSSION

In this study, we successfully reconstituted VSV G-mediated membrane fusion in a minimal model system that enabled time-resolved imaging of single lipid mixing and content release events. The ability to mediated fusion with lipid bilayers at low pH implies that VSV G does not strictly require a proteinaceous receptor to induce fusion. On the other hand, VSVpp fusion was markedly augmented by anionic lipids, POPS, or BMP. Only bilayers containing POPS or BMP supported complete VSVpp fusion, whereas a POPC-based mixture supported only limited lipid mixing activity. Our results
are in full agreement with the previous report that BMP somewhat selectively facilitates VSV G-mediated lipid mixing with liposomes (31). However, in that study, the beneficial effect of BMP was observed in the background of 50% POPS, thus complicating the evaluation of relative roles of these lipids in VSVpp fusion. We found that, although BMP supported more efficient lipid mixing and faster conversion from hemifusion to full fusion than POPS, these differences were modest. In addition, the average time required to complete the release of viral content, which is inversely proportional to the size of a fusion pore (60), was identical for BMP and POPS bilayers. These findings argue against POPS as the specific receptor for VSV and highlight the potential direct role of BMP in entry of this virus.

The BMP content in late endosomes was reported to be as high as 15% of total phospholipids, with the POPC:BMP molar ratio ~3:1 (32) being close to the lipid composition of the SLB used in our study. By contrast, late endosomal membranes contain only 5% of PS (32), which appears to be even lower than the PS content of the plasma membrane (61, 62). Furthermore, because VSV infection is not inhibited by the PS-binding protein annexin V (22), it is unlikely that the incoming VSV encounters PS at the cell surface or during intracellular trafficking. On the other hand, recent evidence suggests a role for BMP in VSV entry from late endosomes (28). The observation that lipid mixing between VSV and endosomes is not attenuated by BMP depletion with specific antibodies (28) suggests that, similarly to dengue virus (30), VSV may undergo hemifusion with early endosomes. It is conceivable that complete fusion can be delayed until VSV enters into late compartments enriched in BMP (28). However, given the evidence that VSV enters from early endosomes (36, 63), further studies are required to test this model.

The relatively low probability of full fusion between VSVpp and SLBs containing BMP or POPS may be due to incorporation of fewer G proteins compared with bona fide VSV (64, 65). In addition, fusion with liposomes appears to occur at the base of a bullet-shaped VSV particle (65), suggesting that the architecture of the virus and perhaps the interactions between G and M proteins regulate the fusion activity. Another explanation for the poor fusion efficiency is that low pH and anionic lipids may not be the only factors required for VSV fusion. Future studies of single VSV fusion with lipid bilayers should distinguish between these possibilities.

Regardless of the lipid composition of SLBs tested in this study, hemifusion always preceded or full fusion. For some particles, the lag time from lipid mixing to the onset of content release reached several minutes. These results agree with previous data on influenza (45), HIV-1 (55) and avian sarcoma and leucosis virus (66) fusion and show that hemifusion is a true and long-lived intermediate stage of VSV G-mediated fusion. Note, however, that for 10% of single influenza virus fusion events with SLB, viral content transfer preceded lipid mixing (45). No explanation for this rather unusual phenotype has been offered by the authors. In addition, our experiments detected transient closing/shrinkage of fusion pores formed by VSV G, a phenomenon known as pore flickering (38, 56, 58, 59).

In conclusion, the SLB model offers a number of important advantages for mechanistic studies of viral fusion at the single particle level. Also, pseudotyping the MLV core with VSV G or other viral glycoproteins provides a universal platform for future mechanistic studies of single virus fusion. The virus labeling and imaging strategies introduced in this study enable detailed studies of the role of lipids at distinct intermediate steps of VSVpp fusion: hemifusion and formation of a small

FIGURE 5. Kinetics of distinct steps of single VSV pseudovirus fusion. A and B, distributions of the delay times from acidification to the onset of lipid mixing (hemifusion, red circles) to the onset of content release (green circles) as well as the lag times between hemifusion and fusion for a given particle (blue circles) for bilayers containing POPS (A) and BMP (B). Only the lag time between lipid and content transfer was significantly different between POPS and BMP (p < 0.045). C, distributions of the duration of mKO release (i.e. time from the onset of mKO signal decrease to reaching the background fluorescence level) that reflect the relative diameter of fusion pores. The total number of full fusion events was 17 for the POPS mixture (red circles) and 14 for the BMP mixture (green circles). No fusion events were observed for the POPC-based mixture lacking the anionic lipids.
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REFERENCES

Lipid Dependence of VSV G-mediated Single Virus Fusion

and Lehmann, M. J. (2009) Visualizing fusion of pseudotyped HIV-1 particles in real time by live cell microscopy. Retrovirology 6, 84
Supplementary Figure and Movie Legends

Supplementary Figure 1. Mean fluorescence intensity (MFI) profiles of VSV G pseudoparticles that failed to undergo hemifusion or fusion at low pH. The relatively stable signals from single particles indicate that photobleaching occurred at a significantly lower (in some cases undetectable) rate than loss of fluorescence due to DiD or mKO release. Two representative traces are shown for each lipid composition: POPC (A-B), POPS (C-D) and BMP (E-F). Note that quenching of the CF-PE fluorescence upon acidification was often associated with concomitant reduction in the mKO signal due to a spectral bleed through.

Supplementary Movie 1. Complete fusion of a VSV G pseudovirus with BMP-containing bilayers. A pseudovirus co-labeled with DiD (red) and NC-mKO (green) is marked by a white arrow (see the corresponding Figure 2C showing a sub-area from this movie). The size of the area shown here is 30×30 µm. DiD signal is in red and NC-mKO signal in green (the CF-PE fluorescence is not shown for clarity). The frame rate is 10 fps.

Supplementary Movie 2. Hemifusion of a VSV G pseudovirus with BMP-containing bilayer. Images of a pseudovirus co-labeled with DiD (red) and NC-mKO (green) correspond to Figure 2D, but show a somewhat larger area (24×24 µm). The CF-PE signal is not shown for clarity. The frame rate is 10 fps.
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Pedro M. Matos, Mariana Marin, Byungwook Ahn, Wilbur Lam, Nuno C. Santos and Gregory B. Melikyan

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