

# **Murine Leukemia Virus Gag Localizes to the Uropod of Migrating Primary Lymphocytes**

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# **ABSTRACT**

**B and CD4**- **T lymphocytes are natural targets of murine leukemia virus (MLV). Migrating lymphocytes adopt a polarized morphology with a trailing edge designated the uropod. Here, we demonstrate that MLV Gag localizes to the uropod in polarized B cells and CD4**- **T cells. The uropod localization of MLV Gag was dependent on plasma membrane (PM) association and multimerization of Gag but independent of the viral glycoprotein Env. Basic residues in MA that are required for MLV Gag recruitment to virological synapses between HEK293 and XC cells were dispensable for uropod localization in migrating B cells. Ultrastructural studies indicated that both wild-type and basic-residue mutant Gag localized to the outer surface of the PM at the uropod. Late-domain mutant virus particles were seen at the uropod in form of budding-arrested intermediates. Finally, uropods mediated contact between MLV-infected B cells and uninfected T cells to form virological synapses. Our results suggest that MLV, not unlike HIV, accumulates at the uropod of primary lymphocytes to facilitate viral spreading through the formation of uropodmediated cell-cell contacts.**

## **IMPORTANCE**

**Viruses have evolved mechanisms to coordinate their assembly and budding with cell polarity to facilitate their spreading. In this study, we demonstrated that the viral determinants for MLV Gag to localize to the uropod in polarized B cells are distinct from the requirements to localize to virological synapses in transformed cell lines. Basic residues in MA that are required for the Gag localization to virological synapses between HEK293 and XC cells are dispensable for Gag localization to the uropod in primary B cells. Rather, plasma membrane association and capsid-driven multimerization of Gag are sufficient to drive MLV Gag to the uropod. MLV-laden uropods also mediate contacts between MLV-infected B cells and uninfected T cells to form virological synapses. Our results indicate that MLV accumulates at the uropod of primary lymphocytes to facilitate viral spreading through the formation of uropod-mediated cell-cell contacts.**

**Retroviral assembly is driven by the viral precursor polyprotein**<br>Gag that consists of matrix (MA), capsid (CA), and nucleocapsid (NC)  $(1-4)$  $(1-4)$  $(1-4)$ . Each domain serves distinct functions during the viral assembly process. The MA domain mediates binding of Gag to the plasma membrane (PM). The CA domain mediates Gag-Gag interaction required to form immature and mature viral particles. CA consists of two domains, an N-terminal domain (CA-NTD) and a C-terminal domain (CA-CTD) [\(5,](#page-12-3) [6\)](#page-12-4). The CA-NTD facilitates the oligomerization into the hexameric and pentameric rings within the capsid structure. CA-CTD dimers and trimers form the contact of neighboring hexamers and pentamers and are critical for Gag oligomerization and particle formation [\(7,](#page-12-5) [8\)](#page-12-6). NC mediates packaging of genomic RNA into the viral core and initiates the oligomerization of Gag. Gag of various retroviruses encodes additional proteins that play important roles in assembly and release. For instance, p12 from murine leukemia virus (MLV) and p6 from the human immunodeficiency virus (HIV) contain late-domain motifs that are required for the viral particle to pinch off from the PM  $(9-13)$  $(9-13)$  $(9-13)$ . The PPPY motif in MLV p12 recruits NEDD4-like E3 ligases to promote virus release via a pathway that is dependent on the vacuolar protein sorting 4 (VPS4)  $(9-11)$  $(9-11)$  $(9-11)$ .

It is generally accepted that the assembly of MLV and HIV predominantly occurs at the PM or its invaginations [\(14](#page-12-12)[–](#page-12-13)[20\)](#page-12-14). Both HIV Gag and MLV Gag have also been observed to associate with late endosomes and multivesicular bodies (MVBs) in HeLa, HEK293, and T cells and macrophages [\(21](#page-12-15)[–](#page-12-16)[25\)](#page-13-0). The endosome/

MVB pathway has further been suggested to play a role in viral trafficking to assembly sites [\(26](#page-13-1)[–](#page-13-2)[28\)](#page-13-3). Gag has also been observed to colocalize with exosomes/microvesicles (EMVs) that are secreted as exosomes [\(29,](#page-13-4) [30\)](#page-13-5). In migrating lymphocytes, the association of Gag proteins with EMV facilitates their polarization, which may further promote the polarization of the cells [\(30\)](#page-13-5). The localization of MLV Gag in B and CD4<sup>+</sup> T cells, physiologically relevant cell types for MLV infection, has not been characterized extensively.

MA is the primary viral determinant responsible for targeting Gag to the PM. It mediates Gag-PM association via a covalently linked myristoyl group at its N terminus and basic charges [\(31](#page-13-6)[–](#page-13-7) [33\)](#page-13-8). Basic charges are thought to interact with acidic phospholipids that are enriched at the inner leaflet of the PM [\(14,](#page-12-12) [34](#page-13-9)[–](#page-13-10)[38\)](#page-13-11). Neutralization of basic residues in the polybasic cluster leads to a

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Received 17 April 2014 Accepted 21 June 2014 Published ahead of print 25 June 2014 Editor: W. I. Sundquist

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relocalization of Gag to intracellular compartments and severely reduces viral release in HeLa cells and HEK293 cells [\(35,](#page-13-12) [36,](#page-13-13) [39](#page-13-14)[–](#page-13-15) [43\)](#page-13-16). However, one such HIV basic-domain mutant, K29/31E, although still targeted to intracellular membranes, assembles and releases relatively efficiently in macrophages and T cells [\(42,](#page-13-15) [44\)](#page-13-17).

Viruses exploit and manipulate existing cellular structures and cell-cell adhesion for the purpose of efficient spreading [\(45](#page-13-18)[–](#page-13-19)[48\)](#page-13-20). Retroviral cell-to-cell transmission has been shown to occur through broad virological synapses or thin filopodial bridges and nanotubes [\(47,](#page-13-19) [49,](#page-13-21) [50\)](#page-13-22). In most cases, the interaction between the viral glycoprotein Env and its receptor leads to the establishment of cell-cell contacts. Consequently, Env and viral receptor accumulate at the cell-cell interface [\(49,](#page-13-21) [51](#page-13-23)[–](#page-13-24)[54\)](#page-13-25). In transformed cell lines, the establishment of adhesion between MLV-infected and uninfected cells is followed by the polarized assembly of particles at the virological synapse [\(51\)](#page-13-23). A single tyrosine residue in the cytoplasmic tail of Env is critically involved in polarizing assembly [\(51,](#page-13-23) [55\)](#page-13-26). Basic residues in matrix domain (MA) and Gag multimerization are then needed for MLV Gag recruitment to the viro-logical synapse [\(56\)](#page-13-27).

Immune cells are the natural host cells for most retroviruses, including MLV, which infects T and B cells [\(52,](#page-13-28) [57\)](#page-13-29). Migrating leukocytes, including T and B lymphocytes, display a polarized morphology. The front extension of a migrating cell is referred to as the leading edge, while the rear protrusion is referred to as the uropod [\(58](#page-13-30)[–](#page-13-31)[61\)](#page-13-32). Uropods are enriched in the adhesion molecules intercellular adhesion molecule 1 (ICAM-1), ICAM-2, and ICAM-3, P-selectin glycoprotein ligand-1 (PSGL-1), CD43 and CD44, cholesterol, and the ganglioside GM1 [\(62](#page-13-33)[–](#page-13-34)[65\)](#page-13-35). The microtubule-organizing center (MTOC) and various organelles, including the endoplasmic reticulum (ER), Golgi apparatus, and mitochondria, locate toward the uropod [\(59,](#page-13-36) [66,](#page-14-0) [67\)](#page-14-1). Uropods are also characterized by actin-rich PM projections such as microspikes and microvilli [\(59,](#page-13-36) [66,](#page-14-0) [68\)](#page-14-2).

The uropod has also been implicated in viral spreading. In polarized HIV-infected T cells, viral Gag and genome localize to the uropod [\(54,](#page-13-25) [69,](#page-14-3) [70\)](#page-14-4). MLV has also been observed in association with the uropod of polarized lymphocytes [\(71\)](#page-14-5). In addition, Gag-laden uropods preferentially mediate contacts between infected and uninfected T cells to form virological synapses [\(54,](#page-13-25) [69\)](#page-14-3). Interference with T cell polarity using myosin inhibitors blocks subsequent HIV cell-to-cell transmission [\(69\)](#page-14-3). Mechanistically, HIV uropod localization requires Gag multimerization primarily mediated by the NC domain [\(69\)](#page-14-3). Gag first associates with PM microdomains, which then laterally migrate to the uropod. Gag localization to these microdomains depends on basic residues within the polybasic region in the Gag MA domain [\(72\)](#page-14-6). The prevailing model for a role of the uropod in HIV transmission is that it serves as a platform for HIV assembly and mediates the formation of virological synapses [\(69,](#page-14-3) [72\)](#page-14-6). Whether MLV assembles and buds at the uropod in polarized murine lymphocytes has not been studied in detail.

Here we demonstrate that MLV Gag localizes to the uropod of polarized B cells and CD4<sup>+</sup> T cells. Unlike the polarization of MLV Gag to virological synapses in transformed cell lines, the recruitment of Gag to the uropod was Env independent. Also, basic residues that are required for localization of Gag to virological synapses in transformed cell lines were dispensable for Gag uropod localization. Rather, PM association and Gag multimerization drove uropod localization of MLV Gag. Ultrastructural

studies demonstrated that MLV particles accumulate at the outer surface of the uropod in polarized B cells. Finally, we provide evidence that uropods mediate contact between MLV-infected B cells and target T cells, leading to the formation of virological synapses. Our results support a model in which MLV, like HIV, accumulates at the uropod to facilitate viral spreading through uropod-mediated cell-cell contacts.

#### **MATERIALS AND METHODS**

**Plasmids and reagents.** Proviral pLRB303 vectors expressing wild-type or mutant Friend MLV (F-MLV) Gag-green fluorescent protein (Gag-GFP), Gag-mCherry, or Gag-cyan fluorescent protein (Gag-CFP) ( $\Delta$ Pol) were previously described [\(55,](#page-13-26) [56\)](#page-13-27). An F-MLV Gag-GFP/ $\Delta$ Env construct was generated by introducing two stop codons in Env. F-MLV Gag-GFP/EnvmCherry was generated by inserting the mCherry sequence into the position of*env* corresponding to amino acid (aa) 273. In the CA-CTD mutant, the C-terminal domain of capsid (from GQYQ to RHRE) has been deleted. Yellow fluorescent protein (YFP) fusion proteins of Lamp1, Rab7, Syt7, and VPS4 [\(73\)](#page-14-7) were introduced into the retrovirus-based vector pLZRS to transduce primary cells. Moloney MLV Gag-YFP was previously described [\(21\)](#page-12-15).

**Virus preparation.** Viruses were generated in HEK293 cells by transfection of plasmid pLRB303 or pLZRS, plasmid encoding F-MLV GagPol (to compensate for the absence of Pol), and F-MLV Env (to increase virus infectivity). Fugene6 (Promega) was used for transfection of HEK293 cells. Cell supernatant was filtered after 24 and 48 h using a  $0.45$ - $\mu$ m-poresize nylon membrane filter and stored at  $-80^{\circ}$ C.

**Primary cell infection.** Isolation and *in vitro* transduction of primary mouse CD4<sup>+</sup> T cells and B cells were described previously [\(52\)](#page-13-28). To remove residual virus, cells were treated with trypsin for 5 min at 37°C 8 to 12 h postspinoculation. Eight hours after trypsin treatment, cells were plated on coverslips coated with  $0.5 \mu g/ml$  of recombinant mouse ICAM-1 (R&D systems) and incubated for 1 to 2 h before fixation with 4% paraformaldehyde (PFA) or coculturing with target cells for live-cell imaging.

**Live-cell imaging.** A polyclonal S49.1 T cell line stably expressing mCAT-1-mCherry was generated and used as target cells. Primary murine B cells expressing Gag-GFP were cocultured with target cells, and the formation of F-MLV synapses was monitored using time-lapse spinning disc confocal video microscopy as described previously [\(51,](#page-13-23) [55,](#page-13-26) [56\)](#page-13-27). The formation of virological synapses was observed 1 h after initiation of cocultures. The images were analyzed using Volocity software (PerkinElmer).

**Immunostaining.** For immunostaining, primary antibodies against the following were used: Lamp1 (ID4B) [\(21\)](#page-12-15), CD43 (S7; BD Pharmingen), CD44 (IM7; BD Pharmingen), or goat anti-mCherry (Biorbyt). Cells were incubated with the primary antibody for 30 min, washed three times with phosphate-buffered saline (PBS)–1% bovine serum albumin (BSA), and then incubated with Alexa Fluor 569- or Alexa Fluor 647-conjugated secondary antibodies. After fixation with 4% PFA, cells were analyzed using a spinning disc confocal microscope. The Gag-uropod polarization index was calculated as the ratio of Gag-GFP mean fluorescence intensity (MFI) at the uropod relative to Gag-GFP MFI outside the uropod. Both MFIs were normalized to background GFP MFI outside cells.

**Electron microscopy.** Correlative fluorescence and scanning electron microscopy (SEM) was performed as previously described [\(49\)](#page-13-21). A diamond pencil was used to mark regions of interest for easy reidentification under the SEM. Transmission electron microscopy (TEM) was performed as previously described [\(49,](#page-13-21) [74\)](#page-14-8).

## **RESULTS**

**MLV Gag localizes to the uropod in polarized primary CD4**- **T and B lymphocytes.** T and B lymphocytes are the natural targets of MLV [\(52,](#page-13-28) [57\)](#page-13-29). To determine the distribution of MLV Gag in



<span id="page-2-0"></span>**FIG 1** MLV Gag localizes to the uropod in polarized primary T and B lymphocytes. (A and B) Primary CD4- T cells (A) and primary B cells (B) were infected with F-MLV Gag-GFP (green). Sixteen to 20 h after infection, cells were plated on ICAM-1-coated coverslips, immunostained for the uropod marker CD43 or CD44 (red), and examined by spinning disc confocal microscopy. Depicted images represent the merged extended-focus view of an entire Z-stack. The mean polarization indices of 10 images that quantify the accumulation of Gag-GFP at the uropod are shown at the upper right corners of the Gag-GFP images. (C) Primary B cells were infected with viruses labeled with Gag-YFP (green) and containing full-length Gag-mCherry genomes (red). Eight to 12 h postinfection, cells were treated with trypsin as indicated and monitored by spinning disc confocal microscopy. Scale bars, 10  $\mu$ m.

polarized T and B cells, we infected primary mouse  $CD4^+$  T and B cells with full-length F-MLV encoding GFP-labeled Gag (Gag-GFP). Following the infection, cells were treated with trypsin to remove residual input virus. Cells were plated on ICAM-1-coated coverslips to induce cell migration and polarization. Cells were fixed at 1 h postplating, and Gag-GFP distribution was investigated by spinning disc confocal microscopy. We found that MLV Gag-GFP accumulated at the uropod in both infected CD4<sup>+</sup> T cells and B cells [\(Fig. 1A](#page-2-0) and [B\)](#page-2-0). In primary CD4<sup>+</sup> T cells, Gag showed a strong colocalization with the uropod markers CD43 and CD44 [\(Fig. 1A\)](#page-2-0). In primary B cells, Gag similarly colocalized with the uropod marker CD44 [\(Fig. 1B\)](#page-2-0). Most splenic B cells do not express CD43 [\(75\)](#page-14-9). To quantify the extent of polarization, we calculated a polarization index defined as the ratio of Gag-GFP mean fluorescence intensity (MFI) found at the uropod relative to Gag-GFP's MFI outside the uropod. We defined polarization as an index value greater than 3 for an analysis of at least 10 cells [\(Fig. 1A](#page-2-0) and [B\)](#page-2-0). The theoretical value for the lack of polarity at the uropod (random distribution) would be 1.

To confirm that the observed Gag signal represented *de novo*synthesized Gag and not the residual input virus, we generated viruses in HEK293 cells by cotransfecting plasmids expressing Gag-YFP independently of other viral factors, with full-length F-

MLV expression plasmids encoding MLV Gag-mCherry. Following infection of B cells, a YFP signal would represent residual input virus, whereas an mCherry signal independent of YFP would indicate *de novo*-synthesized Gag. Without trypsin treatment, only a few YFP-labeled input viruses were observed, some of which localized to the uropod. However, following trypsin treatment, YFP signal was not detected [\(Fig. 1C\)](#page-2-0). In both cases, Gag-mCherry was the predominant signal and localized to the uropod in polarized cells [\(Fig. 1C\)](#page-2-0). Collectively, these results indicate that *de novo*synthesized MLV Gag localizes to the uropod of polarized lymphocytes.

**Env is not required for MLV Gag localization to the uropod.** Since B cells express MLV proteins to higher levels than primary CD4- T cells and are able to form virological synapses*in vivo* [\(52\)](#page-13-28), we concentrated our subsequent studies on primary B cells. To determine if MLV Env also localizes to the uropod, we infected primary B cells with F-MLV encoding both Gag-GFP and EnvmCherry. In infected cells, Env-mCherry was observed at the uropod, where it colocalized with Gag-GFP [\(Fig. 2A\)](#page-3-0). Intracellular Env signal that did not colocalize with Gag could be detected near the base of the uropod, likely representing the trans-Golgi network as well as some ER-associated foci [\(Fig. 2A\)](#page-3-0). In transformed cell lines, MLV Env is essential for the establishment of virological



<span id="page-3-0"></span>**FIG 2** The uropod localization of Gag is independent of Env. (A) Primary B cells were infected with full-length F-MLV encoding Gag-GFP (green) and Env-mCherry (red) and monitored by spinning disc confocal microscopy. The mean polarization index of Env-mCherry or Gag-GFP accumulation at uropods is shown at the upper right corner. (B) Primary B cells infected with F-MLV Gag-GFP (green) lacking envelope ( $\Delta$ Env). The mean polarization index of Gag-GFP accumulation at uropods is shown at the upper right corner. (C) Primary B cells infected with F-MLV Env-mCherry (red) ( $\Delta$ Gag) lacking Gag. The mean polarization index of Env-mCherry accumulation at uropods is shown at the upper right corner. Scale bars, 10  $\mu$ m.

synapses and the subsequent recruitment of Gag to virological synapses [\(51\)](#page-13-23). To test if Env is required for MLV Gag localization to the uropod, we infected primary B cells with F-MLV Gag-GFP/  $\Delta$ Env virus lacking Env. In the absence of Env, Gag still localized to the uropod, indicating that the localization of Gag to the uropod is an intrinsic feature of Gag [\(Fig. 2B\)](#page-3-0). To test if MLV Env localizes to uropod in the absence of Gag, we infected primary B cells with F-MLV Env-mCherry/ $\Delta$ Gag virus lacking Gag. In the absence of Gag, Env still localized to the uropod as well as some intracellular compartments near the base of the uropod [\(Fig. 2C\)](#page-3-0).

**MLV Gag PM binding and multimerization are required for localization to the uropod.** To identify viral determinants within Gag that are responsible for uropod localization, we examined a panel of Gag mutants depicted in [Fig. 3](#page-4-0) [\(55,](#page-13-26) [56\)](#page-13-27). A G2A mutant that abrogates myristoylation of Gag was evenly distributed in the cytoplasm, indicating that myristoylation-mediated membrane binding was required for MLV Gag uropod localization [\(Fig. 3\)](#page-4-0). However, the MA domain was expendable for uropod targeting provided that MA was replaced with a heterologous membrane targeting domain (MTD), in this instance from Src kinase. MLV Gag p12 and NC domains were also not required for uropod targeting [\(Fig. 3\)](#page-4-0), in contrast to CA and CA-CTD mutants that no longer targeted the uropod despite clearly binding to the PM and additional intracellular membranes. The polarization indices of CA and CA-CTD deletion mutants are slightly higher than the theoretical value of 1, presumably because the uropod is a membrane-rich structure and many organelles localize to this area.

Considering the requirement for CA-CTD, we hypothesized that Gag multimerization was critical for Gag uropod targeting. While deletion of either MA or NC alone was insufficient to abolish Gag uropod targeting, deletion of both ( $Src\Delta MA\Delta NC$ ) blocked Gag uropod targeting in the context of an intact CA-CTD domain. The ability of SrcΔMAΔNC to localize to the uropod was restored by introducing a trimeric leucine zipper domain (LZ) at the position of NC that stimulates Gag-Gag interactions independently of Gag-RNA binding [\(56,](#page-13-27) [76\)](#page-14-10). The observation that  $Src\Delta MAANC$  (= Src-p12-CA) does not localize to the uropod reveals that the CA-CTD is necessary but not sufficient for Gag uropod targeting. Taken together, these data support a model in which multiple Gag multimerization signals are involved in regulating Gag targeting to the uropod.

**Basic residues in MA are not required for MLV Gag uropod localization.** We have previously shown that basic residues, irrespective of whether they are contributed by the endogenous polybasic region of MA or come from heterologous MTDs such as Src MTD, are strictly required for the polarized recruitment of Gag to virological synapses in transformed tissue culture cell lines [\(56\)](#page-13-27). This conclusion was based on several previous observations. First, replacing Gag MA with the Lck MTD (Lck $\Delta$ MA) lacking basic residues supported Gag membrane targeting but not recruitment to virological synapses. In contrast, Gag  $\Delta MA$  mutants were targeted efficiently to virological synapses by the Src MTD ( $Src\Delta MA$ ) that contains a myristoylation site and 4 basic residues. Second, a basic patch mutant (bm) of Gag in which the net charge of the polybasic region in the MA was neutralized by replacing two basic residues with acidic residues (K31KRR34 to EKER) assembled intracellularly and did not polarize to the virological synapses in transformed cell lines. Third, N-terminally adding the MTDs of Src but not Lck in front of the MA bm mutant (Srcbm and Lckbm) restored the polarization of MLV to virological synapses [\(56\)](#page-13-27).



<span id="page-4-0"></span>**FIG 3** PM binding and Gag multimerization target MLV Gag to the uropod. The upper scheme displays tested wild-type (wt) MLV Gag and mutants of MLV Gag. All mutations were made in a full-length virus construct with GFP fused to the C terminus of Gag. "LZ" represents the trimeric leucine zipper domain. Localization of indicated Gag-GFP mutants (green) within polarized primary B cells was determined as described for [Fig. 1B.](#page-2-0) Scale bars, 10  $\mu$ m.

To determine if basic residues play a similar role in MLV Gag localization to the uropod, we examined the localization of Src $\Delta$ MA, Lck $\Delta$ MA, Srcbm, Lckbm, and bm Gag-GFP constructs in primary B cells. Src $\Delta$ MA, Lck $\Delta$ MA, Srcbm, and Lckbm Gag-GFP all displayed strong uropod localization, suggesting that MLV Gag uropod localization in primary B cells is independent of the basic residues in MA [\(Fig. 4;](#page-5-0) [Table 1\)](#page-5-1). To our surprise, the bm mutant, which is mislocalized to intracellular compartments in HEK293 cells, showed a clear uropod localization in primary B cells [\(Fig. 4;](#page-5-0) [Table 1\)](#page-5-1). However, the bm mutant appears to be somewhat impaired in membrane binding compared to the wild type. This is visible in its increased cytoplasmic fluorescence and also reflected in the relatively low polarization index, 3.45. Still, Gag uropod targeting in primary lymphocytes is fundamentally different from Gag virological synapse targeting in transformed tissue culture cell lines. Gag uropod targeting requires Gag N-terminal myristoylation and Gag multimerization, in contrast to targeting to virological synapses, which requires these signals in addition to basic residues in Gag's N terminus and upstream Env signaling.

**Wild-type Gag and bm Gag do not localize to late endosomes in primary B cells.** It has been previously reported that neutralization of the basic residues in the polybasic cluster of MLV interferes with PM association and leads to reduced viral release in NIH 3T3 and HEK293 cells [\(36\)](#page-13-13). In these cell lines, the bm mutant localizes to intracellular structures, presumably late endosome/ MVBs. Because the bm mutant was targeted to the uropod in primary B cells, we asked if wild-type and bm Gag in primary B cells might localize to late endosomes in close proximity to the uropod. We first analyzed the distribution of wild-type and bm Gag-GFP F-MLVs in HEK293 cells. Wild-type MLV Gag was targeted to both the PM and late endosomes/MVBs labeled by Lamp1





**SrcAl Lck**<sup>1</sup> Srcb Lckb bn

<span id="page-5-0"></span>**FIG 4** Basic residues in MA are not required for MLV Gag uropod localization. Upper scheme displays tested wild-type (wt) MLV Gag and mutants of MLV Gag. Localization of indicated Gag-GFP mutants (green) within polarized primary B cells was determined as described for [Fig. 1B.](#page-2-0) Scale bars, 10  $\mu$ m.

and Rab7 [\(Fig. 5A\)](#page-6-0), consistent with our previous findings [\(21\)](#page-12-15). bm Gag almost exclusively colocalized with the late endosome markers Lamp1 and Rab7 [\(Fig. 5A\)](#page-6-0), confirming previous reports [\(36\)](#page-13-13). MLV-infected B cells were immunostained with an antibody against Lamp1 or cotransduced to express the late endosomal marker YFP-Rab7. Although Lamp1- and Rab7-positive vesicles

<span id="page-5-1"></span>**TABLE 1** Summary of polarization phenotypes of MLV mutants in virological synapses (VS) in fibroblasts and at uropod

<b>MLV</b>	Polarization		Polarization index (uropod localization) $(n = 10)$	
	VS in fibroblasts <sup>a</sup>	Uropod	Mean	SEM
Wild type (T cells)	$^{+}$	$^{+}$	6.75	2.44
Wild type (B cells)	$^{+}$	$^{+}$	9.63	1.46
$\Delta$ Env		$^{+}$	10.90	1.80
G2A			0.91	0.06
$Src\Delta MA$	$^{+}$	$^{+}$	5.38	1.03
$\Delta$ P12	$^+$	$^{+}$	44.32	5.50
$\Delta$ CA	$^{+}$		1.43	0.05
$\Delta$ CA-CTD	$^{+}$		1.37	0.12
$\Delta NC$	$^{+}$	$^{+}$	3.53	0.56
SrcΔMAΔNC			1.41	0.06
Src $\Delta$ MA $\Delta$ NCLZ	$^{+}$	$^{+}$	7.92	1.75
<b>SrcNC</b>	$^+$		1.39	0.03
Lck∆MA		$^{+}$	8.03	2.28
Srcbm	$^{+}$	$^{+}$	6.56	1.29
Lckbm		$^{+}$	7.42	0.94
bm		$^{+}$	3.45	0.46
$\Delta$ PPPY	$^+$	$^+$	14.18	2.50

*<sup>a</sup>* Determined by Li et al. [\(56\)](#page-13-27). VS, virological synapses.

localized to the base of the uropod, little colocalization was observed with wild-type or bm Gag and either of these markers [\(Fig. 5B\)](#page-6-0). Thus, unlike with the aforementioned cell lines, neither wild-type nor bm Gag localizes to late endosomes/MVBs in primary B cells.

Although neither wild-type nor bm Gag localized to late endosomes/MVBs in primary B cells, it was still possible that they would localize to other intracellular compartments just beneath the PM. To address this issue, we developed an assay based on MLV Env accessibility. The rationale is that if the MLV virions localize intracellularly, Env of the virions should not be accessible to antibodies against the Env ectodomain without permeabilization of the cellular membranes. We tested this assay in HEK293 cells, in which wild-type and bm virions accumulate intracellularly [\(Fig. 6A\)](#page-6-1). Cells were transfected with plasmids encoding F-MLV Gag-GFP/Env-mCherry. After 24 h, cells were fixed and incubated with Alexa Fluor 647-conjugated anti-mCherry antibody and examined by epifluorescence microscopy. The GFP signal and mCherry signal represented the total Gag and total Env, respectively. Indeed, wild-type or bm Gag-GFP/Env-mCherry viruses accumulating in late endosomes/MVB in HEK293 cells were both mCherry and Gag positive but Alexa Fluor 647 negative in the absence of membrane permeabilization [\(Fig. 6A\)](#page-6-1). In contrast, Env-mCherry expressed at the PM and Env-mCherry colocalizing with wild-type Gag-GFP puncta at the PM were accessible to the Alexa Fluor 647-conjugated anti-mCherry antibody without permeabilization [\(Fig. 6A\)](#page-6-1). When we applied this assay to primary B cells, all Env-mCherry signals colocalizing with wild-type or bm Gag-GFP puncta were accessible to the Alexa Fluor 647-labeled anti-mCherry Env antibody without permeabilization, indicating surface accessibility for both wild-type and bm virions [\(Fig. 6B\)](#page-6-1).

**PM of the uropodis a site of active virion production for both wild-type and bm Gag.** The resolution of light microscopy could not determine if the PM of the uropod was the site of active virion



<span id="page-6-0"></span>**FIG 5** Wild-type Gag and bm Gag do not localize to late endosome/MVB in primary B cells. (A) HEK293 cells were transfected with plasmids expressing Lamp1-YFP or YFP-Rab7 (yellow) as well as a plasmid encoding F-MLV Gag-CFP (cyan) and examined 24 h posttransfection by spinning disc confocal microcopy. A single plane of a Z-stack is displayed. (B) Primary B cells were transduced with a viral construct expressing F-MLV Gag-CFP (cyan) and immunostained for Lamp1 (red) or cotransduced with viral constructs expressing YFP-Rab7 (yellow). Images depict an entire Z-stack merged into a single extended-focus view. Scale bars, 10  $\mu$ m.



<span id="page-6-1"></span>**FIG 6** Virions produced by wild-type and bm Gag accumulate at the outer surface of the PM in primary B cells. (A) HEK293 cells were transfected to express wild-type and bm mutant F-MLV Gag-GFP/Env-mCherry. One day later, cells were fixed, incubated with Alexa Fluor 647-conjugated antimCherry antibody, and examined by epifluorescence microscopy. The GFP and mCherry signals represent the total Gag and total Env, respectively. The Alexa Fluor 647 signal represents the surface-exposed Env. (B) An experiment as in panel A was performed with primary B cells. Single Z-planes are displayed. Scale bar, 10  $\mu$ m.

assembly, as opposed to Gag accumulating in an unassembled form at the cytoplasmic face of the plasma membrane. To determine if virion formation occurred at the PM of the uropod, we performed transmission electron microscopy (TEM) of primary B cells expressing wild-type or bm Gag. To preserve a normal morphology of viruses, primary B cells were transduced to express wild-type GagPol in addition to F-MLV Gag-GFP virus. To assess the role of the mutant bm Gag in uropod localization, this mutant was also introduced into the GagPol context of F-MLV. Virusgenerating B cells were fixed and examined by confocal microscopy before being processed for TEM. The addition of GagPol or bm GagPol did not alter the uropod localization of wild-type or bm Gag-GFP, respectively (data not shown). Uropods could be easily identified under TEM based on the polarized morphology of B cells. In polarized B cells, cellular organelles such as the ER, Golgi apparatus, lysosomes, and mitochondria were polarized to-ward the uropod [\(Fig. 7A](#page-7-0) and [B\)](#page-7-0). Importantly, electron-dense virions containing wild-type and bm Gag localized outside the PM and associated thin membrane protrusions or microvilli at the uropod [\(Fig. 7A](#page-7-0) and [B\)](#page-7-0). No intracellular virions were observed in B cells examined, supporting the absence of virus assembly on intracellular compartments like MVBs in primary B cells [\(Fig. 7\)](#page-7-0). For wild-type viruses, on average, 3.5 viral particles were at the uropod and 0.03 viral particle was outside the uropod [\(Fig. 7C\)](#page-7-0). The bm mutant showed a similar distribution [\(Fig. 7C\)](#page-7-0). It is worth noting that TEM thin sections are typically 50 to 70 nm thick, while the uropod is typically 2 to 5  $\mu$ m thick. Consequently, 3.5 uropod-localized viral particles in TEM micrographs equals hundreds of viral particles on the surface of one uropod in threedimensional space.

To investigate the localization of MLV at the uropod in greater detail, we performed correlative fluorescence and scanning electron microscopy (SEM). The samples were prepared as described above, and MLV Gag-GFP-expressing B cells were first identified by fluores-



<span id="page-7-0"></span>**FIG 7** Wild-type and bm viruses localize to the surface of uropods in primary B cells. (A and B) Primary B cells were infected with wild-type (A) or bm (B) full-length F-MLV Gag-GFP viruses in the additional presence of wild-type and bm mutant viruses carrying GagPol. One day after infection, B cells were fixed and processed for TEM. The images to the right are magnified views of regions marked by dashed squares. The inset at the right bottom corners are magnified virus particles from indicated areas. Arrows indicate the positions of virions. Scale bars, 10  $\mu$ m. (C) Quantification of uropod-localized and non-uropodlocalized viral particles for wt Gag, bm Gag, and PPPY Gag in TEM micrographs. Sixty, 27, and 28 polarized B cells with viral particles were analyzed for wt Gag, bm Gag, and  $\overline{\Delta}$ PPPY Gag, respectively. Error bars represent the standard errors of the means.

cence microscopy and then visualized by SEM. Correlated micrographs revealed that uropods of uninfected B cells have abundant thin membrane protrusions or microvilli [\(Fig. 8A\)](#page-8-0). However, infected primary B cells with a strong Gag-GFP signal at the uropod exhibited spherical particles uniform in size  $(\sim 150 \text{ nm})$  and morphology in SEM micrographs for both wild-type and bm Gag [\(Fig. 8B](#page-8-0) and [C\)](#page-8-0). The abundance and the close proximity of the viruses at the uropod

do not permit the identification of single Gag-GFP-labeled virus particles. However, the distribution pattern of spherical  $\sim$ 150-nm structures observed in SEM broadly matched the distribution pattern of Gag-GFP fluorescence, suggesting that these spherical particles are MLV virions [\(Fig. 8B](#page-8-0) and [C\)](#page-8-0). Thus, the SEM data also support the notion that both wild-type and bm virions undergo active assembly and budding at the uropod.



<span id="page-8-0"></span>**FIG 8** SEM electrographs of polarized primary B cells reveal surface accumulation of wild-type and mutant bm Gag particles. Samples were prepared as described for TEM in the legend to [Fig. 7.](#page-7-0) After fixation, virally infected Gag-GFP-labeled (green) polarized B cells were identified using fluorescence microscopy and marked using a diamond pen, and the identical cells were visualized by SEM. Images of uninfected B cells (A) and B cells infected with wild-type (B) and mutant bm Gag (C) F-MLV are displayed. The images to the right are magnifications of indicated regions. The bottom left image of each panel is the corresponding fluorescence image of MLV Gag-GFP virions.

To directly visualize the localization of budding intermediates, we expressed the MLV late-domain mutant ( $\Delta$ PPPY) in B cells. This mutant still localized to the uropod [\(Fig. 9A\)](#page-9-0), similar to the earlier phenotype observed for the p12 deletion that contains the PPPY motif [\(Fig. 3\)](#page-4-0). Both Gag mutants appeared to accumulate at the uropod, resulting in very high polarization indices, consistent with an inability of late-domain mutant virus to pinch off from the membrane. Indeed, TEM revealed that 15.7% of viral particles were still connected with the PM, probably representing budding intermediates of MLV virions [\(Fig. 9B](#page-9-0) and [C\)](#page-9-0). In contrast, very few such structures were observed in samples containing virus expressing wild-type or bm Gag [\(Fig. 9C\)](#page-9-0). Taken together, these data suggested that PM of the uropod is a site of active virus production.

**Uropods mediate contact between MLV-infected B cells and target cells.** Uropod-associated HIV has been proposed to serve as a preformed platform for the formation of virological synapses to facilitate virus transmission to uninfected cells [\(69,](#page-14-3) [72\)](#page-14-6). We performed live-cell imaging to test if uropod-associated MLV also participates in virological synapse formation and MLV cell-to-cell transmission. Primary B cells expressing Gag-GFP labeled virions were cocultured with S49.1 mouse T cells stably expressing mCAT-1-mCherry. The formation of MLV virological synapses was monitored by time-lapse microscopy. Virological synapses



<span id="page-9-0"></span>**FIG 9** Late-domain mutant virus accumulates at the uropod but fails to pinch off from the PM. (A) Primary B cells infected with F-MLV Gag-GFP  $\Delta$ PPPY as described for [Fig. 1B.](#page-2-0) Scale bar,  $10 \mu m$ . (B) Samples were prepared as described for TEM in the legend to [Fig. 7](#page-7-0) to characterize primary B cells infected with late-domain mutant MLV ( $\Delta \Delta$ PPPY Gag). Scale bar, 500 nm. (C) Quantification of assembly intermediates and assembled particles for wt Gag, bm Gag, and  $\Delta$ PPPY Gag in TEM micrographs. Sixty, 27, and 28 polarized B cells were analyzed for wt Gag, bm Gag, and APPPY Gag, respectively.

between infected B cells and target T cells exhibited a strong accumulation of Gag-GFP and mCAT-1-mCherry at the cell-cell interface [\(Fig. 10\)](#page-9-1). We often observed two distinct phenotypes. First, uropod-mediated contact between primary B cells and target T cells was long-lived, maintained for  $>$  30 min [\(Fig. 10\)](#page-9-1). Donor and target cells often remained dynamic, with their leading edges migrating into different directions, while the uropod-mediated contact remained unchanged. Second, due to the migratory dynamics of cells, uropod-mediated contacts could transiently stretch into long filopodium-like contacts [\(Fig. 11\)](#page-10-0).

Uropod-mediated contacts were also observed between infected and uninfected primary B cells in TEM and SEM micrographs [\(Fig. 12\)](#page-11-0). In thin-section images, viral particles were ob-served to localize within or around the narrow synaptic cleft [\(Fig.](#page-11-0) [12A\)](#page-11-0). Thin membrane extensions were abundant around or



<span id="page-9-1"></span>**FIG 10** An example of uropod-mediated stable virological synapses between primary B cells and T cells. Primary B cells infected with wild-type F-MLV Gag-GFP (green) as for [Fig. 1B](#page-2-0) were cocultured with S49.1 T cells stably expressing receptor mCAT-1-mCherry (red) and examined by live-cell microscopy. Three image series with 8-min intervals are shown. "D" marks the donor cell and "T" marks the target cell. Scale bar,  $10 \mu m$ .

within the synaptic cleft. Taken together, these results suggest that uropods can participate in the formation of virological synapses between MLV-infected B cells and target T cells.

#### **DISCUSSION**

Viruses have evolved mechanisms to coordinate their assembly and budding with cell polarity to facilitate their spreading [\(47\)](#page-13-19). Cell polarization can be cell adhesion dependent or independent [\(47\)](#page-13-19). In this study, we demonstrated that the viral determinants for MLV Gag to localize to virological synapses in transformed tissue culture cell lines (cell adhesion, Env receptor dependent) and the uropod in polarized B cells (cell adhesion independent) are different. Basic residues in MA that are required for the Gag localization to virological synapses in transformed cell lines are dispensable for Gag localization to the uropod in primary B cells. Basic residues in MA of HIV Gag have been reported to play a critical role in the association of HIV Gag with uropod-directed microdomains (UDMs) before they comigrate to the uropod [\(72\)](#page-14-6). A similar role may be played by basic residues in MLV MA during the targeting to an putative acidic interface organized by MLV Env at the virological synapses in transformed cell lines [\(56\)](#page-13-27). However, MLV and HIV MA proteins are both dispensable for Gag uropod localization in polarized lymphocytes [\(69\)](#page-14-3). Consequently, basic residues of MA seem to mediate Gag association with specific PM microdomains but are not necessarily required for Gag uropod localization.

The basic patch (bm) mutant of MLV Gag can no longer efficiently bind to the PM and accumulates in late endosomal/MVB



<span id="page-10-0"></span>**FIG 11** Examples of transitions between filopodia and uropod-mediated virological synapses between primary B cells and T cells. Samples were prepared as described in the legend to [Fig. 10.](#page-9-1) "D" marks the donor cell and "T" marks the target cell. Scale bars, 10  $\mu$ m.

compartments in HEK293 cells [\(Fig. 5\)](#page-6-0) [\(36\)](#page-13-13). In contrast, bm Gag is targeted to the uropod and viral particles accumulate at the outer surface. These observations recall previous observations with HIV, for which specific basic mutants were similarly relocalized to late endosomal membranes in HEK293T and HeLa cells but were released in macrophages and primary T cells [\(42,](#page-13-15) [44\)](#page-13-17). Apparently, primary lymphocytes contain alternative trafficking and release pathways that are distinct from those in transformed cell lines.

Gag multimerization is required for both MLV Gag localization to virological synapses in transformed cell lines and Gag uropod localization in polarized B cells. However, MLV NC plays a more important role in Gag localization to virological synapses in transformed cell lines, whereas MLV CA plays a more important role in Gag localization to the uropod. The reason for this difference is unclear.

Previous studies had proposed that any protein can be targeted to the posterior pole, the precursor of uropod, on the basis of PM binding and higher-order oligomerization [\(29,](#page-13-4) [30\)](#page-13-5). The biogenesis of the poster pole primes cells for later signal induced biogenesis of uropod. Our data provide some support for this model, as we observed that Gag derivatives that do not form fluorescent puncta and displayed a diffused distribution on the PM did not localize to the uropod. In contrast, Gag derivatives that multimerize into fluorescent puncta accumulate at the uropod. Late-domain mutant virus also accumulated at uropods and budding arrested particles unable to pinch off from the PM were clearly visible at the uropod. While we cannot exclude surface movement of budding intermediates initiated elsewhere at the PM, these data are consistent with a model in which the uropod represents an active area of virus assembly and release.

MLV Gag localization to the uropod of primary B cells resem-



<span id="page-11-0"></span>**FIG 12** Electron micrographs of uropod-mediated virological synapses between primary B cells. (A and B) Samples were prepared as for [Fig. 7](#page-7-0) and [8](#page-8-0) and examined by TEM (A) and correlative SEM (B). The images to the right in panel A are magnifications of indicated regions. Arrows in panel A point to positions of virions. Insets in panel B are the bright-field and fluorescence images taken by confocal microscopy. "D" marks the donor cell and "T" marks the target cell. The arrow in panel B indicates the position of the uropod-mediated contact.

bles HIV Gag localization to the uropod of primary T cells in many ways. They both are independent of Env expression and represent an intrinsic feature of the retroviral Gag protein [\(69\)](#page-14-3). They both require Gag multimerization involving several domains of Gag. HIV- and MLV-laden uropods both appear to mediate the formation of virological synapses between infected donor cells and uninfected target cells [\(69\)](#page-14-3). However, NC is required for HIV Gag uropod localization, while it is dispensable for MLV Gag uropod localization [\(69\)](#page-14-3). This is unexpected since in HEK293T cells, HIV Gag lacking NC can assemble into somewhat aberrant particles [\(77\)](#page-14-11), while MLV Gag lacking NC does not assemble at all [\(78\)](#page-14-12). Indeed, when we transfected HEK293 cells with our full-length MLV construct expressing  $\Delta NC$  Gag-GFP, no viral particles were observed [\(55\)](#page-13-26). However, when we infected primary B cells with MLV  $\triangle$ NC Gag-GFP, GFP puncta could be readily detected in polarized cells, and they accumulated at the uropod. These observations indicate that the dependence of MLV assembly on NC is cell type specific.

The contribution of MLV-laden uropods to the formation of virological synapses has not been previously addressed. Longterm live-cell imaging studies described in this report have shed some light on this process and further add to the original model proposed by the Ono group [\(69\)](#page-14-3) [\(Fig. 13\)](#page-12-17). If uropods can make the initial contact with a target cell, it will likely engage target cells to form virological synapses, as previously proposed [\(69\)](#page-14-3). In contrast, if the leading edge of a migrating lymphocyte makes the initial contact with a target cell, the leading edge will continue to migrate and bypass the target cells. However, uropods, with their many adhesion proteins and Env-laden virions, adhere to the receptor-expressing target cells, while the leading edge continues to drive cellular polarization of the migrating cells. We observed that uropod-mediated contact in this tug of war between migration and cell-cell adhesion is highly dynamic and can stay in a dynamic equilibrium between long filopodium-like contacts and broad uropod contacts.

In sum, our data provide confirmatory evidence that MLV in

![](_page_12_Figure_1.jpeg)

<span id="page-12-17"></span>**FIG 13** Model for uropod-mediated formation of virological synapses in migrating lymphocytes. The leading edge of the migrating donor cell (green) makes an initial contact with a target cell (red), continues to migrate, and passes the target cell until a uropod-mediated contact is established. Continued migratory forces can lead to long filopodial contacts. "D" depicts the donor cell and "T" the target cell. The green and red thick lines represent the enrichment of Gag-labeled viral particles (green) and viral receptor (red), respectively. The green arrow denotes the direction of migration for donor cells.

addition to HIV can utilize uropods for viral spreading via virological synapses. We report differences with respect to the role of Gag multimerization in the targeting to uropods between both viruses and document the distinct role for basic residues in cell adhesion-induced polarity in virological synapses in transformed cell lines and migration-induced polarity in primary B cells. We also contribute to the characterization of the ultrastructural morphology of polarized lymphocytes infected with MLV and uropod-mediated virological synapses. Live-cell imaging supports the model that uropods participate in the formation of virological synapses between primary cells.

#### **ACKNOWLEDGMENTS**

We thank Christin Herrmann for molecular cloning, Zhenting Jiang and Barry Piekos for assistance with scanning electron microscopy, and Xinran Liu for assistance with transmission electron microscopy.

This research was supported by grant R01 CA098727 from the National Institutes of Health to W.M. and Leopoldina Fellowship LPDS2009-21 to X.S.

We declare that no competing interests exist.

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