Leukocyte-specific protein 1 interacts with DC-SIGN and mediates transport of HIV to the proteasome in dendritic cells

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Dendritic cells (DCs) capture and internalize human immunodeficiency virus (HIV)-1 through C-type lectins, including DC-SIGN. These cells mediate efficient infection of T cells by concentrating the delivery of virus through the infectious synapse, a process dependent on the cytoplasmic domain of DC-SIGN. Here, we identify a cellular protein that binds specifically to the cytoplasmic region of DC-SIGN and directs internalized virus to the proteasome. This cellular protein, leukocyte-specific protein 1 (LSP1), was defined biochemically by immunoprecipitation and matrix-assisted laser desorption/ionization time-of-flight mass spectrometry. LSP1 is an F-actin binding protein involved in leukocyte motility and found on the cytoplasmic surface of the plasma membrane. LSP1 interacted specifically with DC-SIGN and other C-type lectins, but not the inactive mutant DC-SIGNΔ35, which lacks a cytoplasmic domain and shows altered virus transport in DCs. LSP1 diverts HIV-1 to the proteasome. Down-regulation of LSP1 with specific small interfering RNAs in human DCs enhanced HIV-1 transfer to T cells, and bone marrow DCs from lsp1−/− mice also showed an increase in transfer of HIV-1BaL to a human T cell line. Proteasome inhibitors increased retention of viral proteins in lsp1−/− DCs, and substantial colocalization of virus to the proteasome was observed in wild-type compared with LSP1-deficient cells. Collectively, these data suggest that LSP1 protein facilitates virus transport into the proteasome after its interaction with DC-SIGN through its interaction with cytoskeletal proteins.

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Abbreviations used: BMDC, bone marrow-derived DC; LSP1, leukocyte-specific protein 1; mDC, myeloid DC; MDDC, monocyte-derived DC; pDC, plasmacytoid DC; siRNA, small interfering RNA.
the cytoplasmic domain of the DC-SIGN molecule are critical for the internalization of HIV and other viruses (27, 29). Incoming HIV-1 particles in DCs are internalized by various DC-SIGN–dependent and –independent pathways. A fraction of HIV-1 internalized in DCs is degraded immediately in the lysosomes. Some of the virus that escapes degradation is retained in endocytic compartments within the cytoplasm and is either transmitted by recycling to permissive CD4+ lymphocytes or degraded by the proteasome (30, 31). The process by which DC–SIGN internalizes and transfers HIV-1 is thought to be mediated through classical endocytic and recycling pathways (32); however, other cellular proteins involved in this process are unknown. In this work, we describe an actin binding molecule, leukocyte-specific protein 1 (LSP1), which interacts with the cytoplasmic domain of DC-SIGN and affects the transport of HIV-1 through the DC.

RESULTS

The cytoplasmic domain of DC-SIGN required for HIV-1 internalization interacts with LSP1

DC-SIGN, a C-type lectin on immature DCs, mediates rapid internalization of intact HIV and contributes to enhanced infection in trans of target cells that express CD4 and chemokine receptors (24, 27). Raji B cells were first used to analyze the effects of full-length DC-SIGN and DC-SIGN with cytoplasmic domain truncations, DC-SIGNΔ35, lacking both dileucine and tyrosine–based motifs, and DC-SIGNΔ20, missing the dileucine motif. These DC-SIGN forms showed comparable cell surface expression, although the DC-SIGNΔ35 mutant showed slightly lower expression (Fig. 1 A). Also, human myeloid DCs (mDCs) cultured in GM-CSF or freshly isolated mDCs showed comparable surface expression of DC-SIGN compared with control stained cells (Fig. 1 B); however, less HIV-1 transfer to T cells occurs in those cells not cultured in GM-CSF (Fig. 1 C, right). Freshly isolated mDCs had previously been reported to express low levels of DC-SIGN that increase significantly after incubation with IL-4 (33). Compared with full-length DC-SIGN, when Raji B cells expressing DC-SIGN mutants were incubated with HIV-1ADA (CCR5 tropic) or HIV-1IBB (CXCR4 tropic) at 37°C for 2 h, washed, and incubated with A3R5 or MT2 cells, they failed to mediate enhancement of T cell infection (Fig. 1 C). We and others have previously shown that Raji cells cannot be infected by HIV-1, and so all infection in Fig. 1 C is mediated by trans-infection rather than cis-infection (22, 27). The DC-SIGN mutant data further support this conclusion, and direct capture of virus versus internalization in the Raji DC-SIGN and deletion mutants has been shown previously (27). In addition, CXCR4–tropic virus cannot infect mature or immature DCs (22). 20 μg/ml mAbs to DC-SIGN (BD Biosciences) completely inhibited HIV-1 transfer by Raji B cells to MT2 T leukemia cells and partially inhibited transfer mediated by human DCs when incubated for 2 h at 37°C before and during transfer of HIV-1ADA (Fig. 1 D). Collectively, these data suggest a role for DC-SIGN in mediating uptake and transfer of the virus by human DCs.

To determine whether the cytoplasmic domain of DC-SIGN was required for internalization, HIV-1–GFP–labeled virions were incubated with Raji B cell lines expressing WT DC-SIGN or DC-SIGNΔ35. HIV-1–GFP–labeled virions were incubated at 4°C for 30 min to quantify cell surface binding. To assess internalization, cells were incubated with ~780 ng/ml of HIV-1–GFP–labeled virions at 37°C for 2 h
and treated with trypsin for 5 min before being added to HeLa cells expressing CD4/CCR5 (MAGI-CCR5). Raji cells expressing DC-SIGN∆35 showed comparable cell surface expression and were able to bind GFP-labeled HIV-1 virions (Fig. 2 A). It has been shown previously (27) that HIV-1 pseudotypes bind to DC-SIGN and the other DC-SIGN mutants. However, in contrast to cells expressing WT DC-SIGN, cells with DC-SIGN∆35 did not internalize HIV-1 (Fig. 1, B and C, left vs. right panel). Thus, internalization appears to be critical for mediating trans-enhancement of HIV infection.

Because internalization and trans-enhancement of HIV infection are linked and mediated by the cytoplasmic region of DC-SIGN, we sought to identify cellular proteins that interacted with this region. DC-SIGN, DC-SIGN∆35, and DC-SIGN∆20 were immunoprecipitated from whole cell extracts of their respective Raji lines with a DC-SIGN mAb that reacted with the extracellular portion of the molecule. The proteins in the immunoprecipitates were separated by two-dimensional gel electrophoresis and visualized by Coomassie blue staining. Two cellular proteins coprecipitated with WT DC-SIGN but were absent in DC-SIGN∆35 and DC-SIGN∆20 control samples. These proteins, ∼45 and 50 kD, were identified by matrix-assisted laser desorption/ionization time-of-flight mass spectrometry as LSP1 (Fig. 3 A) and actin (not depicted). To confirm the mass spectrometry data, Raji cells expressing DC-SIGN or the nonfunctional deletion mutants DC-SIGN∆35 and DC-SIGN∆20 were lysed and...
Figure 4. DC-SIGN associates with LSP1 through a region distinct from its actin binding domains. (A) The caldesmon-like CI and CII (hatched) and villin-like VI and VII (gray) on LSP1 that bind F-actin are indicated. The early truncation mutants generated by site-directed mutagenesis at amino acids 305, 275, and 180 are shown. (B) 293T cells (10^6 cells/well) in a six-well plate were cotransfected with full-length DC-SIGN and full-length LSP1 or the mutants (1–305, 275, and 180) as indicated. Cell lysates prepared 72 h after transfection were immunoprecipitated with anti–DC-SIGN antibody and assayed for LSP1 (lanes 7–10) and DC-SIGN (lanes 11–14) by immunoblot. Cell lysates of transfected cells were analyzed by immunoblot for LSP1 expression in full-length and mutant 1–305, 1–275, and 1–180 constructs (lanes 15–18). (C) 35S-labeled in vitro–cotranslated full-length LSP1 interacts directly with DC-SIGN. Full-length LSP1 or LSP1(1–180) cDNA were in vitro cotranslated with DC-SIGN in rabbit reticulocytes in the presence of 35S. Labeled proteins were immunoprecipitated in the presence of protease inhibitors with monoclonal anti–DC-SIGN conjugated to agarose beads. Immunoblotting, as seen with the Raji cell line, confirmed the interaction between DC-SIGN and LSP1 (Fig. 3 C). MDDCs express higher levels of DC-SIGN when compared with human mDCs and can be isolated in greater cell numbers.

DC-SIGN binds to a distinct domain of LSP1 between amino acids 276 and 304

LSP1 is an F-actin binding protein involved in cell motility, cell adhesion, and IgM internalization (34–37). The caldesmon-like (CI and CII) and villin-like (VI and VII) are two domains on LSP1 that bind F-actin to facilitate neutrophil motility (38). To map the domain of LSP1 that interacted with DC-SIGN, stop codons (TAG) were introduced at amino acids 180, 275, and 305, generating several LSP1 mutants (Fig. 4 A). LSP1(1–305) lacked the villin-like domains (VI and VII) and LSP1(1–275) contained only the caldesmon-like (CI and CII) domains, whereas LSP1(1–180) lacked both the caldesmon-like (CI and CII) and villin-like (VI and VII) domains. LSP1 and the mutants were transfected with DC-SIGN in 293T cells. 72 h after transfection, cells were harvested, and cell lysates were immunoprecipitated with anti–DC-SIGN antibody. Immunoblots showed that DC-SIGN interacted with WT LSP1 and LSP1(1–305) that lacked villin-like domains but failed to interact with LSP1(1–275) that contained only the caldesmon-like domains and LSP1(1–180) that lacked both the caldesmon-like and villin-like domains. These data show that amino acids 276–304, which are independent of the F-actin binding regions, are essential for interaction between DC-SIGN and LSP1.
LSP1 (Fig. 4 B) and suggest that LSP1 interaction with the cellular cytoskeleton is required for its function. To determine if this interaction was direct or the product of multiple proteins, $^{35}$S-labeled in vitro cotranslated full-length LSP1 or LSP1(1–180) and DC-SIGN proteins were immunoprecipitated in the presence of protease inhibitors with a monoclonal DC-SIGN antibody conjugated to agarose beads and electrophoresed on a 10% polyacrylamide gel. The proteins were visualized by autoradiography, showing a direct interaction between the full-length LSP1, but not the mutant (Fig. 4 C).

To examine the interactions of LSP1 with other C-type lectins, LSP1 and LSP1(1–180) were cotransfected with DC-SIGN, L-SIGN, and Langerin in 293T cells. Cell lysates were immunoprecipitated with the respective C-type lectin antibodies. Immunoblots revealed that all C-type lectins interacted with LSP1 (Fig. 4 D, lanes 1–3) and not LSP1(1–180) (Fig. 4 D, lanes 4–6), suggesting a role for LSP1 in other C-type lectin-dependent processes. To address the binding specificity of C-type lectins’ cytoplasmic domain with LSP1, full-length LSP1 was also incubated with irrelevant molecules (CD2, CD4, and CD40L) and showed interaction with CD2 and CD40L, but not the CD4 molecule (Fig. 4 E).

**LSP1 down-modulation by specific small interfering RNAs (siRNAs) in human DCs facilitates HIV-1 transfer to T cells**

To determine whether LSP1 was detectable in immature and mature mDCs, mDCs from healthy individuals were examined before and after maturation with poly:IC. Both immature and mature mDCs expressed LSP1, and the levels did not change with maturation (Fig. 5 A), even though HIV-1 transfers more efficiently in mature mDCs (22). The physiological consequences of LSP1–DC-SIGN interactions were further studied by LSP1-specific siRNAs. Four LSP1-specific siRNAs (Fig. 5 B) were synthesized, and their effectiveness in down-regulating LSP1 was determined initially in Raji cells. Two siRNAs effectively decreased endogenous LSP1 in Raji B cells (Fig. 5 B, siRNAs in A and C), and this knockdown was most efficient at 24 h, with LSP1 levels returning to normal in 48 h. Down-regulation of LSP1 did not change expression of DC-SIGN, CD80, or MHC class II; however, it enhanced the transfer of HIV-1 to T cells (Fig. 5 A, available at http://www.jem.org/cgi/content/full/jem.20061604/DC1).

To determine the effect of HIV-1 internalization and transfer to T cells, human mDCs (incubated with poly:IC), MDDCs, and plasmacytoid DCs (pDCs) isolated from HIV-1 donors were transfected with the GeneSilencer reagent containing siRNAs for LSP1 (siC or siA) or a negative control (siRNA scramble). In transfected cells, siRNAs were mixed with an unrelated fluorescent Cy5-labeled scrambled siRNA at a ratio of 4:1 and identified by flow cytometry after 8 h. Cells were sorted for the Cy5 label, plated onto a 96-well tissue culture plate, and pulsed 24 h after transfection with a single round replication-competent virus expressing a luciferase reporter (HIV-1$_{ADA}$) for 2 h at 37°C, washed, and incubated with A3R5 (CCR5-tropic) T leukemia cells. Cells were assayed for luciferase activity 72 h after transduction. The results demonstrate that LSP1 siRNA-treated Raji cells and various populations of human DCs facilitated greater...
transfer of HIV-1 to T cells compared with control (Fig. 5, C and D). These data suggest that LSP1 interacts with DC-SIGN and/or other C-type lectins to direct the virus away from pathways mediating trans-infection.

**Increased HIV-1 transfer to human T cells by lsp1<sup>+/−</sup> DCs**

To investigate the role of LSP1 in HIV-1 trafficking through DCs, we first determined whether DCs from LSP1 knockout mice conferred the same enhancement of infection to T cells observed in human DCs. To test this hypothesis, bone marrow–derived DCs (BMDCs) were isolated from lsp1<sup>+/+</sup> and lsp1<sup>−/−</sup> mice (34). DCs were incubated with CpG oligonucleotide for 24–48 h to induce maturation. Total protein from immature and mature BMDCs was assayed for LSP1 expression. As with human DCs, expression levels were not affected by maturation (Fig. 6 A), and lsp1<sup>−/−</sup> DCs were null for LSP1 (Fig. 6 B). To determine the effect of LSP1 in murine DCs, mature BMDCs from lsp1<sup>+/+</sup> and lsp1<sup>−/−</sup> mice were incubated with live HIV-1 BaL (~780 ng/ml) for 2 h at 37°C, washed extensively, and incubated alone or with A3R5 T cells. p24 levels were assayed in supernatants 72 h after infection. Similar to knockdown in human DCs, murine DCs that lacked LSP1 increased transfer of virus to T cells (Fig. 6 C). Because the murine DCs mimic the effect seen in human DCs, they were used to further understand the mechanisms of LSP1-mediated HIV-1 internalization and the infection of T cells.

**DC LSP1 traffics HIV-1 to the proteasome for degradation**

Incoming HIV-1 particles in DCs are bound and internalized by various DC-SIGN–dependent and –independent pathways. Immediately after internalization, most of the virions are degraded in an acidic lysosomal compartment. A fraction of the virus that escapes degradation is retained in endocytic compartments and is either transmitted to permissive CD4<sup>+</sup> lymphocytes or degraded by the proteasome (30, 31). To investigate the role of LSP1 in DC processing and transport of HIV-1, HIV-1–GFP was incubated with DCs in the presence or absence of proteasome or lysosome inhibitors. Because of the inability to confirm the complete silencing of LSP1 and the limited window of knockdown in human mDCs, the LSP1 knockout mice provide the best model system to study the degradative pathway in the total absence of LSP1. Differentiated BMDCs were incubated with the lysosomal inhibitors chloroquine (100 μM), the endosome acidification inhibitor bafilomycin A1 (10 μg/ml), or the proteasomal inhibitor MG132 in RPMI (5 μg/ml) or RPMI alone for 1 h before transduction with concentrated HIV-1–GFP for 2 h. Cells were washed extensively and incubated with the respective inhibitors. Before lysis, cells were treated with trypsin and washed to remove virus bound to the cell surface. Total protein isolated at various time points after the 2-h transduction (0 h, 1 h, and 3 h) was assayed by ELISA for p24 Gag. Results at 1 h showed that more retention of HIV-1 occurred in lsp1<sup>−/−</sup> BMDCs when compared with lsp1<sup>+/+</sup> controls (Fig. 7 A, left). The p24 values at other times showed a similar trend as the 1-h time point, but the effect was maximal at 1 h. In lsp1<sup>+/+</sup> control cells treated with the proteasome inhibitor MG132, HIV-1 retention was higher compared with chloroquine–, bafilomycin–, or vehicle-treated lsp1<sup>+/+</sup> cells (Fig. 7, middle) and were similar to lsp1<sup>+/−</sup> BMDCs when treated with the same inhibitors (Fig. 7, right), suggesting that LSP1 may be involved in the shuttling of virus to the proteasome for degradation.

To determine the colocalization of HIV-1 to the proteasome in DCs, mature BMDCs from lsp1<sup>−/−</sup> and wt mice were pulsed with HIV-1–GFP for 30 min at 37°C. For proteasome inhibitor studies, BMDCs from lsp1<sup>−/−</sup> and wt mice were incubated with the proteasomal inhibitor MG132 in RPMI (5 μg/ml) or RPMI alone for 1 h before infection by concentrated HIV-1–GFP for 30 min. Cells were washed extensively and stained with a mAb to the 20S proteasome subunit a-4. BMDCs from lsp1<sup>−/−</sup> mice showed more HIV-1 retention and less colocalization to the proteasome (Fig. 7 B). In contrast, those cells from wt mice showed greater colocalization of HIV-1 to the proteasome (Fig. 7 B, yellow). Once DCs were treated with the proteasome inhibitor MG132, colocalization was markedly diminished in both lsp1<sup>−/−</sup> and

![Figure 6. Murine DCs from lsp1<sup>−/−</sup> mice increase HIV-1 uptake and transfer to T cells. (A) LSP1 expression in murine DCs remains constant after maturation. Cell lysates prepared from immature or CpG-matured bone marrow–derived murine DCs were assayed for LSP1 expression by a polyclonal anti-LSP1 antibody. (B) LSP1 expression in mature BMDCs isolated from lsp1<sup>+/−</sup>, lsp1<sup>−/−</sup>, and lsp1<sup>+/+</sup> littermates. Cell lysates prepared from CpG-matured bone marrow–derived murine DCs were assayed for LSP1 expression by a polyclonal anti-LSP1 antibody. (C) LSP1-null BMDCs enhance HIV-1 trans-infection of T cells. Mature BMDCs (10<sup>5</sup>/well) from wt (wt) or lsp1<sup>−/−</sup> mice were pulsed with HIV-1<sub>1BaL</sub> (~780 ng/ml) of p24/ml) or RPMI alone for 2 h at 37°C. Cells were washed extensively to remove free virus and replaced with fresh media, and A3R5 T cells were added to one set of mock or HIV-1–infected BMDCs followed by incubation for another 72 h. At the appropriate time, cell supernatants were collected, and p24 ELISA was performed as instructed by the manufacturer (Coulter). The data are representative of duplicate experiments.](https://example.com/fig6.png)
Down-regulation of LSP1 siRNAs in human DCs or wt mice (Fig. 7 C). Collectively, these results are consistent with the model that LSP1 shuttles the internalized pool of HIV-1 to the proteasome and that lack of LSP1 facilitates HIV-1 transfer.

**DISCUSSION**

In this study, we have demonstrated that DC-SIGN, a C-type lectin on DCs that mediates HIV-1 uptake and transfer to T cells, interacts with LSP1, an actin-binding cytoskeletal protein. HIV-1 uptake is dependent on the cytoplasmic domain of the DC-SIGN molecule (Fig. 1). By using immunoprecipitation for biochemical purification and identification by mass spectrometry, we showed that LSP1 interacts specifically with HIV-1. Our results also showed that LSP1 interacts with other C-type lectins, L-SIGN and Langerin, which are present on the human DC. Because these C-type lectins have been shown to mediate HIV-1 transfer independently of DC-SIGN (39), LSP1 may be involved with these C-type lectins’ interaction with HIV-1. Langerin, unlike DC-SIGN, contains a proline-rich region that is likely responsible for binding and rapid internalization of pathogens. CD2 and CD40L both contain this proline-rich region, possibly providing the region necessary for interaction with LSP1. Down-regulation of LSP1 siRNAs in human DCs or murine *lsp1*/−/− BMDCs showed a dramatic increase in the amount of virus transferred from the DC to the susceptible T cell, as LSP1 was shown to facilitate proteasomal degradation of HIV-1.

A leukocytic protein, LSP1 (also known as WP34, pp52, and leufadin) is a 52-kD F-actin binding phosphoprotein expressed in all human leukocytes and leukocytic cell lines (40–42). The basic C-terminal domain contains amino acid sequences homologous to two known F-actin binding proteins, caldesmon and the villin headpiece (36, 37). Although LSP1 is an F-actin binding protein, it is also a very important regulator of microfilamentous cytoskeleton dynamics (34). After HIV-1 uptake in the DCs, it is internalized into a specialized viral endosome, which is distinct of early and late endosomal vesicles (36), where a fraction of virus remains undigested and polarizes to the infectious synapse between the targeted T cells (24, 27, 28). HIV-1 virus that does not polarize is subjected to lysosomal processing and MHC II antigen presentation, or it is degraded by the proteasome (30, 31). Because LSP1 interacts specifically with full-length DC-SIGN and not a truncated cytoplasmic domain mutant, this finding suggests that it is involved with trafficking HIV through the DC. LSP1 has proven important in polarizing the actin cytoskeleton and aiding in motility of the cell. Using proteasome inhibitors and confocal microscopy, we show that LSP1 helps to shuttle the HIV-1 virus into the proteasome, promoting its degradation, a process independent of its interaction with DC-SIGN. In the absence of LSP1, HIV-1 degradation decreases and more virus is able to recycle to the surface, promoting transfer to T cells. We do not know why the proteasomes in murine DCs were not susceptible to bafilomycin as they are in human DCs. Because they are...
isolated differently from the human cells and are grown in cytokines, it is possibly a difference in the patterns of gene expression in these cells, although we cannot exclude a species effect. Proteasomal inhibitors can affect ubiquitin levels in the cells, which could explain the decrease in HIV-1 degradation; however, in the absence of LSP1, there was less co-localization of HIV-1 to the proteasome and no significant difference in HIV-1 degradation in the *lsp1* c-/* BMDCs compared with wt cells treated with proteasomal inhibitors, suggesting that this effect was independent of ubiquitin effects. This experiment confirmed the role of LSP1 in this degradative process.

Our study reveals new insights into HIV-1 trafficking through DCs leading to the enhancement of T cell infection by DC-SIGN—internalized virus. The role of DC-SIGN in trans-infection is not completely understood, in part because other C-type lectins may be involved in the process in some DC populations, and blocking of DC-SIGN with mAbs does not always completely inhibit HIV-1 transfer. Although one previous study contradicted the report of Kwon et al. (27) that point mutants in the tyrosine and dileucine motifs of the cytoplasmic domain of DC-SIGN do not affect gp120 binding, it is important to recognize that the latter study analyzed internalization with gp120 protein rather than virus (44), and the significance of this assay for virus internalization and transfer is questionable. Nonetheless, consistent with the present work, a mutant with a deletion of the cytoplasmic domain in that study showed the same loss of function as seen here examined by transmission of the lentiviral vector.

The discovery that LSP1, an actin-binding molecule, interacts with DC-SIGN has implications for understanding the trans-enhancement of T cell infection by DCs, possibly leading to ways of blocking transfer. Sequestering actin and the cytoskeleton may lead to decreased transfer of HIV-1, but not without possible serious effects on DC viability. Antigen internalized by DCs has been shown to lead to classical MHC II processing, peptide loading, and surface presentation (45). We find that increased transfer of HIV-1 to T cells in the absence of LSP1 is due to decreased HIV-1 degradation in the proteasome; however, insights into the effect of LSP1 on peptide processing and antigen presentation have yet to be investigated. Collectively, these data suggest a role for LSP1 trafficking of HIV-1 to the proteasome for viral degradation. Continued elucidation of HIV-1 trafficking in DCs provides us with a greater understanding of how C-type lectins, such as DC-SIGN, mediate viral uptake and transfer to susceptible target cells.

**MATERIALS AND METHODS**

**Plasmids and siRNA.** The DC-SIGN CITE-GFP and DC SIGNΔ15 CITE-GFP were expressed under a CMV promotor/enhancer. The cDNA-encoding LSP1 (accession no. BI911034) was expressed in pCMV-SPORT6 (Invitrogen). Mutagenesis of LSP1 was performed using a Stratagen Quick Change Site-Directed Mutagenesis kit according to the manufacturer’s directions. To introduce a stop codon to generate LSP1c (1–305, 275, and 180) mutants, the following primers were used: LSP1c (1–305): 5′ primer: GGGAGGCTCCAAGACCTCATATACTGATCAACAT-

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**Virus production, entry, transduction, and infection assays.** Pseudotyped HIV-1ΔΔAΔGFP lentivirus—expressing luciferase was infected by transfection cotransfection of 293T cells using calcium phosphate (Promega). In brief, the packaging vector pMD 8.2, pHR-luciferase, and the envelope expressing p58II-HIVΔΔAΔGFP or pK85-HIVΔΔAΔGFP were transiently transfected into 293T cells. Supernatants were harvested 48 and 72 h after transfection, filtered, and stored at −80°C. Virus concentration was determined by an ELISA assay for the p24 antigen (Beckman Coulter) (16–22).

**GFP-Vpr-labeled HIV-1 lentivirus (HIV-1ΔΔGFP)** was produced by transfection of 293 T cells with the pLAI provirus and the plasmid pEGFP-C3 (CLONTECH Laboratories, Inc.) containing the entire Vpr coding region fused to the carboxyterminus of eGFP (GFP-Vpr). Cells were washed at 16–20 h after transfection and replenished with fresh media. 48 h later, supernatants were harvested, filtered through a 0.45-μm syringe filter, and concentrated. In brief, 32 ml of supernatant was layered on 5 ml of Optiprep (Iodoxinal) medium (Invitrogen) and centrifuged at 50,000 g for 1.5 h with a Surespin 630 rotor (Sorvall). The last 3 ml of supernatant remaining above the Optiprep interface was collected and frozen at −80°C in 300-μl aliquots (27, 29). Concentrated HIV-1ΔΔAΔGFP (MOI, ~1; 7.8 μg/ml p24) was prepared in PBMCs and provided by J. Mascola and M. Louder (Vaccine Research Center, NIAID, NIH).

24 h after transfection of siRNAs, HIV-1ΔΔAΔGFP infection was performed in a 96-well flat-bottomed luciferase plate by removing 200 μl RPMI culture media and adding 200 μl of virus stock. After incubation for 2 h at 37°C, cells were washed twice and incubated with 1.5 × 10^6 A3R5 T cells for 48 h, lysed, and assayed for luciferase activity with a commercially available kit (Promega). Similarly, BMDCs from LSP1 transgenic mice were infected with WT HIV-1ΔΔAΔGFP (MOI, ~1; 7.8 μg/ml p24) for 2 h and washed five times with RPMI. All animal experiments were reviewed and approved by the Animal Care and Use Committee, Vaccine Research Center (VRC), NIAID, and performed in accordance with all relevant federal and NIH guidelines and regulations. The DCs were incubated with A3R5 T cells for 48 h and assayed for p24.

**Lsp1ΔΔ** and C57BL/6 wt mouse BMDCs were isolated and incubated in 20 ng/ml RPMI plus GM-CSF in a 96-well tissue culture dish (5 × 10^5). The cells were matured in 5 μg/ml of ODN CpG (1829) for 24–48 h. Cells were pretreated with 10 μg/ml bafilomycin, 5 μg/ml MG132, 100 μM chloroquine, or alone in RPMI for 1 h. Media were then removed, and DCs were pulsed with HIV-1ΔΔGFP for 2 h. Cells were washed once with PBS and lysed in 1X p24 cell lysis buffer at time points 0 h, 1 h, and 3 h and stored at −30°C. Lysates were assayed for p24 activity by ELISA (Beckman Coulter HIV-1 p24 Antigen EIA) and read at 450/570 nm dual wavelength on a SPECTRAMax Plus 384 (16–22).

**Cells and transfections.** Parental control Raji-1 cells and Raji-1 cells stably transfected with human DC-SIGN (Raji DC-SIGN) or DC-SIGN with a cytoplasmic truncation that lacks both the dileucine motif and the tyrosine-based motif in the cytoplasmic tail (Raji DC-SIGNΔ35 and DC-SIGNΔ20) were provided by D. Littman (New York University School of Medicine, New York, NY) and maintained in RPMI media at 37°C and 5% CO2. A3R5 T cells were provided by J. Mascola (Vaccine Research Center, NIAID, NIH) and cultured in RPMI and gentamicin (G418).
Human mDCs and pDCs were purified from elutrated monocytes from healthy adult donors by a two-step procedure consisting of automated leukapheresis and counterflow centrifugal elutriation at the Transfusion Medicine Department of the Warren Grant Magnuson Clinical Center, NIH. Because the cells only are used and are without identifiers, this work with the cells is exempt from IRB review. mDCs and pDCs were isolated from the elutrated monocyte fraction with negative selection by removing cells expressing BDCA-4 and CD19 with microbeads (Miltenyi Biotec), followed by positive selection using antibodies to CD1c (Miltenyi Biotec). mDCs and pDCs were then cultured in a medium containing 10 ng/ml GM-CSF (PeproTech) for 18–24 h before any experiments. mDCs were also induced to mature using 50 μg/ml poly:IC (Sigma-Aldrich) for 48 h. Human MDCDS were generated from monocytic cells cultured in RPMI, hGM-CSF, and IL-4 for 7 d before phenotyping cells for CD1c1 to ensure purity.

BMDCs were matured using 50 ng/ml ODN CpG (1829) for 24–48 h. All animal experiments were approved by IACUC and conducted in compliance with all relevant federal and NIH policies.

Mature human DCS were transfected with a control of LSP1 siRNA (QiAGEN). For each transfection, 225 nM siRNA (180 nM unlabeled siRNA and 45 nM Cy5-labeled siRNA) was used. Transfections were performed in a 4:1 mixture of unlabeled siRNA and Cy5. To transfect the mature human DCS with siRNA, 10^6 cells were resuspended in 1 ml of serum-free RPMI. The siRNA mixture was combined with 9 μl of GeneSilencer (Genlantis) and added to the cell suspension according to the manufacturer’s protocol. The cells were washed 8 h after transfection and sorted by a fluorescence-activated cell sorter (Becton Dickinson) to identify Cy5-labeled siRNA-transfected cells. Sorted cells were seeded in a 96-well plate at 5 × 10^4 cells/well.

**Immunoprecipitation, immunoblot, and mass spectrometry.** Raji cells expressing DC-SIGN or a nonfunctional mutant, DC-SIGNΔ35 or DC-SIGNΔ20, were lysed in 1X cell lysis buffer (Cell Signaling). 1 or 2 ng of total protein was incubated with monoclonal anti–DC-SIGN antibody for 2 h. Protein G (Invitrogen) conjugated to agarose was then added and incubated overnight at 4°C. The mixture was washed three times with 1X cell lysis buffer and loaded onto a polyacrylamide 4–12% gel for 1-D or 2-D electrophoresis. Gels were stained with Coomassie blue or proteins were transferred to a PVDF membrane (Invitrogen) and assayed for LSP1 and DC-SIGN by immunoblot using polyclonal antibodies.

Protein identification of 1-D or 2-D gel-separated proteins was performed on reduced and alkylated, trypsin-digested samples prepared by standard mass spectrometry protocols. Tryptic digests were chromatographed (1 μl/min), and peptides were separated using a Zorbax C18SBW reverse phase column (0.15 mm ID × 100 mm). The mobile phase consisted of a gradient prepared from solvent A (0.2% formic acid) and solvent B (99.8% acetonitrile, 0.2% formic acid) at room temperature. For these fractionated digests, capillary LC-tandem MS (LC-MS/MS) was performed with a CapLC and a quadruple-time of flight mass spectrometer (Qtof-2; Waters Micromass). Computer-controlled data-dependent automated switching to MS/MS provided peptide sequence information. MassLynx and Global Server software were used for data acquisition and processing. Data processing and databank searching were performed with Mascot software (Matrix Science). The NCBI�r protein database from The National Center for Biotechnology Information, NLM/NIH, was used for the search analysis.

35S-labeled in vitro transcription translation. Full-length LSP1 or LSP1(1–180) cDNAs were in vitro cotranslated with DC-SIGN in rabbit reticulocytes in the presence of 35S (TNT-sportc; Promega). Labeled proteins were immunoprecipitated in the presence of protease inhibitors with monoclonal anti–DC-SIGN conjugated to agarose beads electrophoresed on a 10% polyacrylamide gel. The gel was fixed and dried using a slab gel dryer (Savant SGD 2000) at 80°C for 5 h. Labeled proteins were visualized by autoradiography after 9 h.

**Confocal microscopy.** 10^4 Raji B cells expressing full-length DC-SIGN and cytoplasmic 35 amino acid–deleted DC-SIGNΔ35 were incubated with 100 μl of a GFP-labeled HIV-1 pseudovirus for 2 h. Cells were treated with trypsin-EDTA, washed, and added to MAGI-CCKR5 HeLa cells (10^4 cells/well) plated onto eight-well coverslip slides (Nunc). Sequential images of live cells were recorded every 3 min by confocal microscopy (SP2-AOBS; Leica Microsystems), and uptake, polarization, and transfer were assessed with representative cells.

10^4 mature BMDCs from lpl^-/- and wt mice were pulsed with 100 μl HIV–1–GFP for 30 min at 37°C. For proteasome inhibitor studies, BMDCs from lpl^-/- and wt mice were incubated with the proteasomal inhibitor MG132 in 5 μg/ml RPMI or RPMI alone for 1 h before infection by concentrated HIV–1–GFP for 30 min. Cells were washed once with PBS plus 2% FCS, fixed, permeabilized (BD Biosciences), and stained with a mAb (15 μg/ml) to the 20S proteasome subunit a-4 (BioMol). Cells were viewed using confocal microscopy and analyzed using Leica software.

**Online supplemental material.** Fig. S1 demonstrates that LSP1 down-regulation causes increased HIV-1 transfer in Raji B cells that does not affect normal surface receptor expression. Fig. S1 is available at http://www.jem.org/cgi/content/full/jem.20061604/DC1.

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