plasmid construction, protein expression, crystallization, and structure determination are provided in the Supplemental Data.

Protein-Protein Interaction Experiments

Biosensor binding experiments were performed as described (Garrus et al., 2001) with purified HIV-1 $p6^{Gag}$, EIAV $p9^{Gag}$, and HIV-1 TSG101_{UEV} proteins binding to immobilized GST-ALIX_{Bro1-V}, GST- $\mbox{ALIX}_{\mbox{\scriptsize V}},$ GST-ALIX $_{\mbox{\scriptsize Bro1}},$ and GST-ALIX $_{\mbox{\scriptsize 714-723}}$ proteins. Assay conditions and binding affinities are provided in Table S2. GST pulldown experiments were performed as described (von Schwedler et al., 2003) using purified ALIX_{Bro1} proteins binding to GST or GST-CHMP4A proteins captured from clarified E. coli lysates.

Assays for HIV-1 APTAP Release and Infectivity

293T cells (\sim 8 × 10 5 cells/well in 6-well plates) were transfected with 1 μg of HIV-1 ΔPTAP plasmid (Garrus et al., 2001) + 1 μg of ALIX expression vector per well (10 μl Lipofectamine 2000, Invitrogen). Cytoplasmic proteins and sucrose-pelleted virions were harvested 24 hr posttransfection, analyzed by western blotting, and quantified using an Odyssey imaging system (Li-COR, Inc.). Primary antibodies were rabbit anti-CA and rabbit anti-MA at 1:15,000. HIV-infectious titers were assayed in single-cycle MAGIC assays in P4 cells. Additional experimental details are provided in von Schwedler et al. (2003).

EIAV VLP Production and ALIX Silencing

HeLa M cells (\sim 4 × 10⁵ cells/well, 6-well plate) were transiently transfected (10 µl; FuGene6, Roche Applied Science) with 5 µg of wild-type or p9^{Gag} ΔΥΡ pEV53B EIAV vector (Olsen, 1998). EIAV virus-like particles were harvested 48 hr posttransfection, concentrated through 20% sucrose cushions, and analyzed by western blotting using affinity-purified EIAV anti-CA antibody (1:10,000). For ALIX-depletion experiments, an shRNA targeting human ALIX nucleotides 1765-1783 (Chen et al., 2005) was delivered using the FG12 lentiviral expression vector (Qin et al., 2003; 8 µg/ml polybrene, moi = 15) 24 hr prior to transfection with pEV53B.

Supplemental Data

Supplemental Data include Experimental Procedures, References, five figures, and two tables and can be found with this article online at http://www.cell.com/cgi/content/full/128/5/841/DC1/.

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REFERENCES

Alroy, I., Tuvia, S., Greener, T., Gordon, D., Barr, H.M., Taglicht, D., Mandil-Levin, R., Ben-Avraham, D., Konforty, D., Nir, A., et al. (2005). The trans-Golgi network-associated human ubiquitin-protein ligase POSH is essential for HIV type 1 production. Proc. Natl. Acad. Sci. USA 102, 1478-1483.

Bibollet-Ruche, F., Bailes, E., Gao, F., Pourrut, X., Barlow, K.L., Clewley, J.P., Mwenda, J.M., Langat, D.K., Chege, G.K., McClure, H.M., et al. (2004). New simian immunodeficiency virus infecting De Brazza's monkeys (Cercopithecus neglectus): evidence for a cercopithecus monkey virus clade. J. Virol. 78, 7748-7762.

Bieniasz, P.D. (2006). Late budding domains and host proteins in enveloped virus release. Virology 344, 55-63.

Cabezas, A., Bache, K.G., Brech, A., and Stenmark, H. (2005). Alix regulates cortical actin and the spatial distribution of endosomes. J. Cell Sci. 118, 2625-2635.

Chatellard-Causse, C., Blot, B., Cristina, N., Torch, S., Missotten, M., and Sadoul, R. (2002). Alix (ALG-2-interacting protein X), a protein involved in apoptosis, binds to endophilins and induces cytoplasmic vacuolization. J. Biol. Chem. 277, 29108-29115.

Chen, C., Vincent, O., Jin, J., Weisz, O.A., and Montelaro, R.C. (2005). Functions of early (AP-2) and late (AIP1/ALIX) endocytic proteins in equine infectious anemia virus budding. J. Biol. Chem. 280, 40474-

Dejournett, R.E., Kobayashi, R., Pan, S., Wu, C., Etkin, L.D., Clark, R.B., Bogler, O., and Kuang, J. (2006). Phosphorylation of the proline-rich domain of XP95 modulates XP95 interaction with partner proteins. Biochem. J. 401, 521-531.

Demirov, D.G., and Freed, E.O. (2004). Retrovirus budding. Virus Res. 106.87-102.

Demirov, D.G., Orenstein, J.M., and Freed, E.O. (2002). The late domain of human immunodeficiency virus type 1 p6 promotes virus release in a cell type-dependent manner. J. Virol. 76, 105-117.

Gallop, J.L., and McMahon, H.T. (2005). BAR domains and membrane curvature: bringing your curves to the BAR. Biochem. Soc. Symp. 72,

Garrus, J.E., von Schwedler, U.K., Pornillos, O.W., Morham, S.G., Zavitz, K.H., Wang, H.E., Wettstein, D.A., Stray, K.M., Cote, M., Rich, R.L., et al. (2001). Tsg101 and the vacuolar protein sorting pathway are essential for HIV-1 budding. Cell 107, 55-65.

Geminard, C., De Gassart, A., Blanc, L., and Vidal, M. (2004). Degradation of AP2 during reticulocyte maturation enhances binding of hsc70 and Alix to a common site on TFR for sorting into exosomes. Traffic 5,

Gottlinger, H.G., Dorfman, T., Sodroski, J.G., and Haseltine, W.A. (1991). Effect of mutations affecting the p6 gag protein on human immunodeficiency virus particle release. Proc. Natl. Acad. Sci. USA 88,

Gouet, P., Courcelle, E., Stuart, D.I., and Metoz, F. (1999). ESPript: analysis of multiple sequence alignments in PostScript. Bioinformatics 15, 305-308.

Huang, M., Orenstein, J.M., Martin, M.A., and Freed, E.O. (1995). p6Gag is required for particle production from full-length human immunodeficiency virus type 1 molecular clones expressing protease. J. Virol. 69, 6810-6818.

Hurley, J.H., and Emr, S.D. (2006). The ESCRT complexes: structure and mechanism of a membrane-trafficking network. Annu. Rev. Biophys. Biomol. Struct. 35, 277-298.

Ichioka, F., Horii, M., Katoh, K., Terasawa, Y., Shibata, H., and Maki, M. (2005). Identification of Rab GTPase-activating protein-like protein (RabGAPLP) as a novel Alix/AIP1-interacting protein. Biosci. Biotechnol. Biochem. 69, 861-865.

Katoh, K., Shibata, H., Suzuki, H., Nara, A., Ishidoh, K., Kominami, E., Yoshimori, T., and Maki, M. (2003). The ALG-2-interacting protein Alix associates with CHMP4b, a human homologue of yeast Snf7 that is involved in multivesicular body sorting. J. Biol. Chem. 278, 39104-39113.