Viruses are champions when it comes to commandeering the normal biological processes of their hosts. In the case of viruses with small genomes, such as the retroviral human immunodeficiency virus (HIV) and influenza, the limited coding capacity of the viral genome forces the virus to use many host cell factors to extend its capabilities during entry into, replication within, and budding from the cells of its host. There is currently a lot of interest in understanding how retroviruses interact with their hosts. Work in this area not only helps us understand how viruses replicate, but also sheds light on normal cellular processes.

A great deal of attention has been given to how viruses latch onto cell-surface receptors and hijack other components to enter and replicate in the host cells, but much less is known about how they usurp the cellular machinery to orchestrate their exodus. A recent study has uncovered a new player in this process.

Margaret Wang, Wankee Kim, Pietro De Camilli, Stephen Goff and colleagues have identified a new endocytic protein involved in the production of a retrovirus, the Moloney murine leukemia virus (Mo-MuLV). The authors characterize the association of a protein involved in the formation of endocytic vesicles, endophilin 2, with the Mo-MuLV protein Gag, and a possible role for this interaction in the production of virus particles. The study identifies a potentially significant new player in retrovirus release and opens up a new line of investigation aimed at understanding the interplay between endocytosis and the cellular release of retroviral particles.

All retroviruses have three key genes, encoding proteins called Gag, Env and Pol. Gag is the structural protein that makes up the viral core and drives viral assembly and release. Gag is a polyprotein and is organized into four distinct regions: the matrix (MA) domain, which is closely associated with the plasma membrane and implicated in budding functions; the capsid (CA), which condenses to form ordered core particles; the nucleocapsid (NC), an RNA-binding protein; and a cleavage product whose name, as well as function, varies depending on the virus (for example, for HIV it is called p6 and for Mo-MuLV it is p12).

Late in the infection cycle of Mo-MuLV (see the ‘Background’ box), the viral Gag polyprotein captures the RNA genome, binds to the plasma membrane and assembles into spherical enveloped particles that bud from the cell. Gag is known as the particle-making machine because it can assemble and bud in the absence of other viral proteins. Hence, any
additional machinery necessary for viral budding and membrane fission must be supplied by the cell and recruited by Gag.

**Assays in vitro**

To identify further which cellular factors are recruited by Mo-MuLV during the production and budding of virus particles, Wang et al. [1] used a yeast two-hybrid assay of a mouse T-lymphoma cDNA library using Gag as bait and identified endophilin 2 as a Gag-binding partner. A second yeast two-hybrid screen showed that endophilin 2 interacts specifically with the MA portion of Gag. In vitro binding assays further confirmed the endophilin-Gag association.

Members of the endophilin family of proteins are involved in endocytic vesicle formation. Endophilin 2 is one of three members of the subgroup endophilin A and is a regulatory component of the machinery involved in clathrin-mediated endocytosis. These proteins are known to promote membrane curvature and bending and are involved in vesicular trafficking events of endocytosis [2].

Many suspected that endophilins were going to be involved in budding because of their known function,” says Stephen Goff (from the Department of Microbiology at Columbia University in New York City, USA, and senior author of the Journal of Biology article). “Sometimes, when you do these screens you recover these proteins that are obscure. When we saw endophilin, we realized we had one that made sense.” (See the ‘Behind the scenes’ box for further discussion of the motivation for the work.)

**Towards viral therapeutics**

The question of whether endophilin 2 is actually required for virion production remains somewhat hazy. Wang et al. thought that ‘knocking down’ endophilin 2 levels with a small interfering RNA (siRNA) might cause virus

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**The bottom line**

- According to a current model of enveloped RNA virus egress from cells, the viruses hijack several components of the host endocytic pathway in order to escape the cell. The retroviral Gag protein is the key player in the process of virion assembly at the plasma membrane and is thought to associate with endocytic proteins to promote virion budding.

- A yeast two-hybrid protein-protein interaction screen has identified an endocytic protein, endophilin 2, that interacts with the Gag protein of Moloney murine leukemia virus (Mo-MuLV).

- Endophilins are known to promote membrane curvature and bending and are involved in vesicular trafficking events of endocytosis.

- Endophilin 2 is packaged inside the virion and associates with the matrix portion of Gag.

- Perturbation of endophilin levels affects virion production. Thus, it seems likely that endophilins are also involved in the budding of Mo-MuLV.

- This study increases our understanding of how the endosomal sorting machinery is linked to retrovirus release and will stimulate further investigations aimed at more fully defining the cellular components involved in virus budding.
production to go down. Despite reducing the levels of endophilin 2 by up to 50%, they saw no effect on virion numbers - but the method could not fully eliminate endophilin 2, so the residual protein might have been sufficient for function. In a second attempt at eliminating the wild-type function of endophilin 2, dominant-negative effectors (produced by overexpressing wild-type or fragmented endophilin 2) were used to block the protein’s function, and this did significantly reduce virion production, although it was not eliminated entirely.

These experiments provide evidence that the Gag-endophilin 2 interaction is of functional importance, since viral particle production was inhibited, says Sundquist. “But there is the outstanding question that is raised but not answered by Goff’s work which is why the virus interacts with endophilins. So, we now need to understand why this interaction takes place, and such experiments can be very challenging,” he says.

Heinrich Göttlinger, in the Department of Cancer, Immunology and AIDS at the Dana-Farber Cancer Institute in Boston, USA, adds that “although the work is suggestive, Wang et al. have not yet formally demonstrated that endophilins are indeed required for efficient virus release. Future siRNA knock-downs targeting all the endophilins simultaneously should definitively resolve this issue.”

But Goff says that such an experiment is nearly impossible since it would almost certainly be detrimental to the cells’ survival. Despite the negative results with the knockdown experiment, Goff is not discouraged. “This [negative result] is probably due to redundancy, where other members of the endophilin family compensate for the loss of endophilin 2 expression, or because the remaining levels of endophilin 2 inside the cells are abundant enough to carry out their function.” Indeed, the levels of endophilin 2 needed are likely to be very low, as only a minute amount of the intracellular protein is incorporated into virions [1].

“IT would also be interesting to know whether the reduced levels of virus production induced by endophilin over-expression are due to a block in virus release or whether the effect might be elicited earlier in the virus assembly/release pathway,” says Eric Freed, Chief of the Virus-Cell Interaction Section at the National Cancer Institute in Frederick, USA.

Interestingly, Wang et al. found that the Gag-endophilin association was not conserved among other retroviruses. Using the yeast two-hybrid assay, they tested Gag proteins from HIV, Rous sarcoma virus (RSV), Mason-Pfizer monkey virus (MPMV) and simian immunodeficiency virus (SIV). Endophilin 2 interacted with Gag from RSV, which is closely related to Mo-MuLV, but not with any of the other Gags from more distantly related viruses. According to Goff, endophilin could nevertheless be involved in HIV production, although indirectly. The alternative is that some other protein performs the equivalent function for HIV. “This is entirely possible. But we do not know what that protein or proteins might be,” he adds.

Although it is not yet fully clear how endophilin 2 helps the virus function, the authors do speculate that perhaps endophilin, which normally helps to generate membrane curvature,
Behind the scenes

*Journal of Biology* asked Stephen Goff to comment on the background to the project to search for new host proteins that interact with the Moloney murine leukemia retrovirus.

**What motivated you to look for host cellular proteins that interact with Mo-MuLV?**

We think there are a large number of host cellular proteins that are yet to be appreciated as important for retrovirus replication. Some of these, like endophilin, will likely be involved in one or more of the many steps of virion assembly. Different retroviruses, and viruses of other families, will probably use different subsets of these proteins. There will also probably be even more proteins involved at other stages of the life cycle: viral entry, uncoating, reverse transcription, entry into the nucleus, proviral formation and viral gene expression. Discovering these host factors will not only lead to new antivirals, but also to a greater understanding of cell biology.

**How long did it take your group to do the experiments and what were the steps that ensured success?**

This project has a long history. It actually started eight years ago when Sandy Morse at the NIH did a large-scale screen for cellular proteins that interacted with the Gag protein of a murine leukemia virus. We were doing similar screens with Moloney, but it wasn’t until early 2000 when Margaret Wang, a new graduate student, decided to look over Sandy’s hits again. Margaret discovered that one of his initially unknown genes, recovered many times, was by then identified as endophilin 2. The function of the endophilins made this gene immediately exciting as it could be involved in virus assembly, so she decided to work these hits up. The work described in the paper constituted the major effort of her graduate studies.

**What was your initial reaction to the results and how were they received by others?**

We were excited instantly upon the realization that these hits were an endophilin. It made perfect sense. I remember a conversation with Eric Freed about the general factors that would be exciting to find as involved with virion budding, and he proposed that finding a role for endophilins would be an obvious expectation even before our findings were known. Given the strong evidence that has come out recently that so many host proteins are involved in virion budding and release, I think most people in the field are happy to include them in the list of players.

**What are the next steps?**

There will be some easy work to survey the viruses that are utilizing this machinery, and to survey which factors are needed by each virus. There will be some mapping of the binding between the viral and host proteins. There may be structural studies, including co-crystallization. The next difficult issue for all the proteins recently identified as cofactors is to determine their actual functions in the process. This might involve in vitro virion assembly reactions, with the inclusion of membranes in these reactions. But it is also likely to require a more detailed understanding of membrane biochemistry and behavior.

could be helping the virus to distort the membrane during the process of virion budding. It might also be that the endocytic proteins play an important role in Gag localization and trafficking, as there are now several indications that viral Gag proteins are trafficked via endosomal pathways [3].

Despite the need for more work to clarify some details, Goff’s findings do fit nicely with the general idea that is developing in the field that retroviruses make extensive use of the endosomal machinery to bud from cells [4-6]. For example, the very late stages of HIV-1 release appear to be primarily promoted by a direct interaction between the p6 domain of its Gag and the cellular protein Tsg101, which then connects Gag with the endosomal sorting machinery that is involved in inward budding of vesicles into the multivesicular body (reviewed in [7]).

What’s more, many other proteins that are part of endocytic complexes, such as Vps28, Hrs, AIP-1, Vps4, and a group of proteins called CHMPs, have been shown to associate with Gag via small motifs called L domains, and are required for various very late steps in virion budding and release [7]. Others have recently shown that infectious HIV particles can bud internally into late endosomal compartments in macrophages [8], and that MuLV RNAs (and apparently also Gag proteins) are transported to the cell membrane by endosomal vesicles [3].

“I think that there is a general feeling in the field that all of these observations will ultimately be unified to explain how retroviruses are trafficked along the endosomal system and ultimately released from cells,” says Sundquist. “Another important aspect of Goff’s work is that although it has been clear that retroviruses utilize cellular proteins that function late in the endosomal pathway (for example, Tsg101, AIP1, and Vps4), it has been much less clear that they utilize proteins that function earlier in endocytosis, such as endophilin 2 and clathrin,” he adds.
This new model of virus budding has many implications for both viral pathogenesis and cell biology. Virus budding is an attractive new drug target, because the pathway appears to be essential and common to many viruses - although common targets can also have potential problems arising from their generality. Cellular targets might also be less susceptible to drug resistance than is the case for many conventional viral targets. "It may be possible to inhibit retrovirus release by targeting the lipid-modifying activity of endophilins with small molecule inhibitors," suggests Göttlinger.

Apart from helping design new antiviral drugs, these studies on viral budding will certainly extend our understanding of the endosomal sorting machinery. In many cases, viruses have helped in the discovery of fundamental aspects of cellular biology and facilitated their analysis. For instance, oncoproteins were first identified as host sequences acquired by transforming retroviruses; reverse transcriptase was first identified as a viral enzyme; regulation of translation was first seen in the interferon response to virus infection; and such processes as inhibition of translation and use of internal ribosome entry sites were first studied with poliovirus. Learning how Mo-MuLV and other retroviruses use the endocytic pathway is sure to lead to more new and exciting discoveries for cell biologists.

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