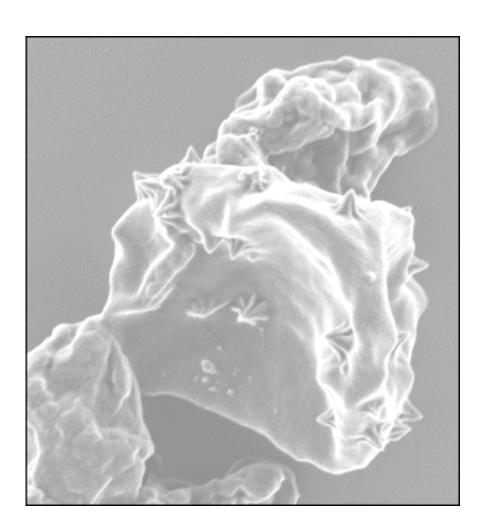


From hot springs to HIV, same protein complexes are hijacked to promote viruses

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This is a scanning electron micrograph of *Sulfolobus solfataricus* cells infected with STIV, showing pyramid-like structures on the surface of the cell. Credit: Montana State University



Biologists from Indiana University and Montana State University have discovered a striking connection between viruses such as HIV and Ebola and viruses that infect organisms called archaea that grow in volcanic hot springs. Despite the huge difference in environments and a 2 billion year evolutionary time span between archaea and humans, the viruses hijack the same set of proteins to break out of infected cells.

In eukaryotes—the group that includes plants and animals—and in archaea—micro-organisms with no defined nucleus in their cellular construction—<u>viruses</u> co-opt a group of important protein complexes called the Endosomal Sorting Complexes Required for Transport, or ESCRT.

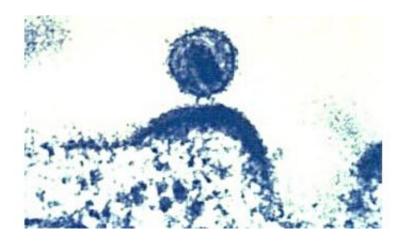
The researchers were studying *Sulfolobus* turreted icosahedral virus, or STIV, which infects *Sulfolobus solfataricus*, a species of archaea called a thermophile that can be found in volcanic springs, such those in Yellowstone National Park. Thermophiles are micro-organisms that survive in extremely hot environments. The researchers found that, as with a range of viruses that infect humans, STIV is also dependent upon its host's ESCRT machinery to complete its life cycle.

"The new work yields insight into the evolution of the relationship between hosts and viruses and, more importantly, presents us with a new and simple model system to study how viruses can hijack and utilize cellular machineries," said Stephen D. Bell, professor in the IU Department of Molecular and Cellular Biochemistry and Department of Biology. Bell is co-lead author on the paper that appears today in early online editions of the *Proceedings of the National Academy of Sciences*.

The researchers looked for interactions between STIV and ESCRT proteins by using a technique in molecular biology called two-hybrid screening, which identifies binding interactions between two proteins or a protein and a DNA molecule. After finding two examples where viral



proteins (the major <u>capsid protein</u> B345 and the <u>viral protein</u> C92) interacted with ESCRT proteins (SSO0619 and SSO0910), epiflouresence microscopy and transmission electron microscopy were used to determine exactly where ESCRT protein components localized in STIV-infected cells.



HIV is shown budding from an infected cell. Similar versions of HIV infect other nonhuman species, such as feline immunodeficiency virus in cats and simian immunodeficiency virus in monkeys and other nonhuman primates. Credit: National Institute of Allergy and Infectious Diseases

Epiflouresence microscopy uncovered spots of the ESCRT protein Vps4 in STIV-infected *S. solfataricus* cells, while no Vps4 was found after similar analysis in uninfected cells. In testing with transmission electron microscopy, the researchers identified Vps4 localized in the seven-sided pyramid-like structures that form in the membrane of *S. solfataricus* prior to viruses causing cell breakdown when the viral protein C92 expressed. No localization of Vps4 was found in similar cells where C92 was repressed.

The work shows that Vps4 is recruited to viral budding sites—those



seven-sided pyramid-like structures—in the *S. solfataricus* thermophile. Significantly, other scientists have shown that the Vps4 protein of the eukaryotic ESCRT machinery localizes to the HIV budding site in humans.

"We believe the ESCRT machinery plays two roles in STIV biology. First, by virtue of interaction between the viral B345 protein and the host protein SSO0619, ESCRT aids in the construction of the STIV viral particles," Bell said. "Second, the strong association we find between the pyramid structures formed by C92 and ESCRT's Vps4 protein allows us to hypothesize that the ESCRT machinery plays a vital role in opening those pyramid exit structures that then leads to cell disruption and the release of viral progeny."

Just as the ESCRT machinery in <u>plants and animals</u> plays a key role in cell division, Bell's lab has previously shown that the same is true for that similar yet less-complicated ESCRT complex in archaea. Also of importance, Bell added, is that the ESCRT apparatus both in eukaryotes and in archaea like *S. solfataricus* is co-opted by viruses.

"These parallels support the idea that the cellular ESCRT is ancient and that it is likely to have evolved prior to archaea and eukarya separating to become different domains of life," Bell said.

Scientists date <u>archaea</u> back to 3.7 billion years, while the oldest eukaryote fossils date back to 1.7 billion years.

More information: "Functional interplay between a virus and the ESCRT machinery in Archaea," June 10, 2013, in *Proceedings of the National Academy of Sciences*: www.pnas.org/cgi/doi/10.1073/pnas.1301605110



Provided by Indiana University

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