

New study shows how the Crimean-Congo hemorrhagic fever virus enters our cells

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Binding of CCHFV glycoproteins to LDLR induces receptor-mediated endocytosis. Credit: *Nature Microbiology* (2024). DOI: 10.1038/s41564-024-01672-3



Researchers at Karolinska Institutet, in collaboration with JLP Health and others, have identified how the tick-borne Crimean-Congo hemorrhagic fever virus enters our cells. The results are <u>published</u> in *Nature Microbiology* and are an important step in the development of drugs against the deadly disease.

Crimean-Congo <u>hemorrhagic fever virus</u> (CCHF virus) is spread through tick bites and can cause hemorrhagic fever.

The disease is serious and has a mortality rate of up to 40 percent, depending on the health status of the person infected.

Common symptoms include fever, muscle pain, <u>abdominal pain</u>, joint <u>pain</u>, vomiting and hemorrhaging that can cause organ failure.

The disease has spread to Europe

The virus is present in around 40 countries, including Central Asia, the Middle East and parts of Africa. In recent years, the disease has spread to new geographical areas as a result of climate change, including Spain and France.

The tick species that can spread the disease has also been observed in Germany and Sweden. There are currently no effective treatments for the disease.

In a new study, researchers at Karolinska Institutet in Sweden and others have found that the virus enters our cells via a protein on the <u>cell surface</u>, the so-called LDL receptors that regulate blood cholesterol levels.

To identify the protein, the researchers used human mini-organs grown



in test tubes and an advanced stem cell library from JLP Health. The same platform has previously been used to identify how the Ebola virus enters cells.

The results were also confirmed in tests on mice, which showed that mice lacking the LDL receptor did not get as sick as others.

Researchers want to trick the virus

The discovery is an important step towards developing drugs for Crimean-Congo hemorrhagic fever, according to Ali Mirazimi, adjunct professor at the Department of Laboratory Medicine, Karolinska Institutet, and one of the researchers behind the study.

"Once we know which receptor the virus uses, we can produce the receptor in test tubes and administer it as a drug," he says. "Then we can trick the virus into binding to those receptors instead of to the cells and thus stop the virus from spreading in our bodies."

This knowledge is essential if the disease were to become more common and spread to new areas. Normally it takes many years to develop a drug, but the COVID-19 pandemic and the development of the SARS-CoV-2 vaccine showed that it can be done much faster if everyone decides it is a priority.

"This is an important step in our preparedness for the disease," says Ali Mirazimi. "Crimean-Congo hemorrhagic fever is a disease we would rather not have. Migratory birds spread the ticks and have already been found in Sweden. If the disease starts appearing in more places, we may already have a drug that we can take into clinical trials."

More information: Vanessa M. Monteil et al, Crimean-Congo Hemorrhagic Fever virus uses LDLR to bind and enter host cells, *Nature*



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Provided by Karolinska Institutet

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