

## **Inhibiting targets of SARS-CoV-2 proteases can block infection, study shows**

September 21 2021





Novel Coronavirus SARS-CoV-2 Transmission electron micrograph of SARS-CoV-2 virus particles, isolated from a patient. Image captured and colorenhanced at the NIAID Integrated Research Facility (IRF) in Fort Detrick, Maryland. Credit: National Institute of Allergy and Infectious Diseases, NIH

Researchers at the University of Liverpool have shown how SARS-CoV-2 viral proteases attack the host cell, and how this can be targeted to stop virus replication in cell culture using existing drugs.

The new findings, published today in *Nature Communications*, offer a powerful resource to understand proteolysis in the context of viral infection, and to inform the development of targeted strategies to inhibit the virus that causes COVID-19.

SARS-CoV-2 has been responsible for over 227 million infections, and more than 4.6 million deaths worldwide during the pandemic. Efforts to test, treat and vaccinate against the virus all benefit from an improved understanding of the basic biology of SARS-CoV-2.

Both viral and cellular proteases play a crucial role in SARS-CoV-2 replication, and inhibitors targeting proteases have already shown success at inhibiting SARS-CoV-2 in cell culture models.

In this study, led by the University of Liverpool and the Institut Pasteur in Paris, researchers used a mass spectrometry approach to study proteolytic cleavage events during SARS-CoV-2 infection.

"Mass spectrometry-based approaches to identify <u>protease</u> substrates have existed for a number of years however, they have seen only limited application to the study of viral substrates, and had not been previously applied to the study of proteolysis during coronavirus infection,"



explains lead author Dr. Emmott Edward, a Tenure-Track Fellow at the University's Institute of Systems, Molecular and Integrative Biology.

The team found previously unknown cleavage sites in multiple <u>viral</u> <u>proteins</u>, including major antigenic proteins S and N, which are the main targets for vaccine and antibody testing efforts.

They discovered significant increases in cellular cleavage events consistent with cleavage by SARS-CoV-2 main protease (Mpro) and identified 14 potential high-confidence substrates of the main and papain-like proteases, validating a subset with *in vitro* assays.

They went on to show that siRNA depletion of these cellular proteins inhibits SARS-CoV-2 replication, and that drugs targeting two of these proteins: the tyrosine kinase SRC and Ser/Thr kinase MYLK, showed a dose-dependent reduction in SARS CoV-2 titres.

Both Bafetinib (an experimental cancer drug) and Sorafenib (an approved drug used to treat kidney and liver cancer) showed SARS-CoV-2 inhibition at concentrations that did not result in cytotoxicity in a human cell line model of infection.

Dr. Emmott said: "An improved understanding of the exact ways in which proteolytic cleavage is regulated, modulates protein activity, and serves to benefit viral replication will be crucial for targeting cellular substrates of viral proteases as a therapeutic strategy.

"As further SARS-CoV-2 variants emerge, the incorporation of post translational modification data from studies such as this can also support efforts to predict phenotypes from genetic data on emerging variants."

**More information:** Characterising proteolysis during SARS-CoV-2 infection identifies viral cleavage sites and cellular targets with



therapeutic potential, *Nature Communications* (2021). DOI: 10.1038/s41467-021-25796-w

## Provided by University of Liverpool

Citation: Inhibiting targets of SARS-CoV-2 proteases can block infection, study shows (2021, September 21) retrieved 28 April 2024 from <u>https://phys.org/news/2021-09-inhibiting-sars-cov-proteases-block-infection.html</u>

This document is subject to copyright. Apart from any fair dealing for the purpose of private study or research, no part may be reproduced without the written permission. The content is provided for information purposes only.