

Blood test could predict who is likely to get long COVID

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BLOOD TEST COULD PREDICT WHO IS LIKELY TO GET LONG COVID

156 Healthcare workers were tested for COVID-19

35% tested +ve

Of those infected, 20% had at least one persistent symptom one year after infection

Proteins in the blood were monitored from 1 week prior to and 6 weeks after confirmed infection, and compared to those who were not infected

Results show that COVID-19 infection leads to a dramatic difference in levels of some of the proteins up to six weeks following infection

Using an artificial intelligence (AI) algorithm, a "signature" in the abundance of different proteins was identified that predicted persistent symptoms a year after infection

What does this mean? A test could potentially be offered alongside a PCR test that could predict people's likelihood of developing Long Covid *

- Results need to now be replicated in a large, independent sample

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[https://www.thelancet.com/journals/ebiom/article/PIIS2352-3964\(22\)00475-3/fulltext](https://www.thelancet.com/journals/ebiom/article/PIIS2352-3964(22)00475-3/fulltext)

Blood test could predict who is likely to get long COVID. Credit: UCL

A blood test taken at the time of COVID-19 infection could predict who is most likely to develop long COVID, suggests a new small-scale study

led by UCL researchers.

The study, published in *eBioMedicine*, analyzed proteins in the blood of health care workers infected with SARS-CoV-2, comparing them to samples from health care workers who had not been infected.

Usually [protein levels](#) in the body are stable, but the researchers found a dramatic difference in levels of some of the proteins up to six weeks following infection, suggesting disruption to a number of important biological processes.

Using an artificial intelligence (AI) algorithm, they identified a "signature" in the abundance of different proteins that successfully predicted whether or not the person would go on to report persistent symptoms a year after infection.

The researchers say that if these findings are repeated in a larger, independent group of patients, a test that could predict people's likelihood of developing long COVID could potentially be offered alongside a [polymerase chain reaction](#) (PCR) test.

Lead author Dr. Gaby Captur (MRC Unit for Lifelong Health and Ageing at UCL) said, "Our study shows that even mild or asymptomatic COVID-19 disrupts the profile of proteins in our blood plasma. This means that even mild COVID-19 affects normal biological processes in a dramatic way, up to at least six weeks after infection.

"Our tool predicting long COVID still needs to be validated in an independent, larger group of patients. However, using our approach, a test that predicts long COVID at the time of initial infection could be rolled out quickly and in a cost-effective way. The method of analysis we used is readily available in hospitals and is high-throughput, meaning it can analyze thousands of samples in an afternoon."

Senior author Dr. Wendy Heywood (UCL Great Ormond Street Institute of Child Health and Great Ormond Street Hospital) said, "If we can identify people who are likely to develop long COVID, this opens the door to trialing treatments such as antivirals at this earlier, initial infection stage, to see if it can reduce the risk of later long COVID."

For the study, researchers analyzed blood plasma samples from 54 health care workers who had PCR- or antibody-confirmed infection, taken every week for six weeks in spring 2020, comparing them to samples taken over the same period from 102 health care workers who were not infected.

They used targeted [mass spectrometry](#), a form of analysis that is extremely sensitive to tiny changes in the amount of proteins in [blood plasma](#), to look at how COVID-19 affected these proteins over the course of six weeks.

The researchers found abnormally high levels of 12 proteins out of the 91 studied among those infected by SARS-CoV-2, and that the degree of abnormality tracked with the severity of symptoms.

The research team found that at the time of first infection, abnormal levels of 20 proteins studied were predictive of persistent symptoms after one year. Most of these proteins were linked to anti-coagulant (anti-clotting) and anti-inflammatory processes.

A machine learning algorithm, trained on the [protein](#) profiles of the participants, was able to distinguish all of the 11 health care workers who reported at least one persistent symptom at one year from infected health care workers who did not report persistent symptoms after a year. Another machine learning tool was used to estimate the likelihood of error and suggested a possible error rate of 6% for this method.

The study was carried out at the UCL Biological Mass Spectrometry

Centre (BMSC).

More information: Plasma proteomic signature predicts who will get persistent symptoms following SARS-CoV-2 infection, *EBioMedicine* (2022). [DOI: 10.1016/j.ebiom.2022.104293](https://doi.org/10.1016/j.ebiom.2022.104293)

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