

A serological assay to detect SARS-CoV-2 seroconversion in humans

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Abstract

SARS-Cov-2 (severe acute respiratory disease coronavirus 2), which causes Coronavirus Disease 2019 (COVID19) was first detected in China in late 2019 and has since then caused a global pandemic. While molecular assays to directly detect the viral genetic material are available for the diagnosis of acute infection, we currently lack serological assays suitable to specifically detect SARS-CoV-2 antibodies. Here we describe serological enzyme-linked immunosorbent assays (ELISA) that we developed using recombinant antigens derived from the spike protein of SARS-CoV-2. Using negative control samples representing pre-COVID 19 background immunity in the general adult population as well as samples from COVID19 patients, we demonstrate that these assays are sensitive and specific, allowing for screening and identification of COVID19 seroconverters using human plasma/serum as early as two days post COVID19 symptoms onset. Importantly, these assays do not require handling of infectious virus, can be adjusted to detect different antibody types and are amendable to scaling. Such serological assays are of critical importance to determine seroprevalence in a given population, define previous exposure and identify highly reactive human donors for the generation of convalescent serum as therapeutic. Sensitive and specific identification of coronavirus SARS-Cov-2 antibody titers may, in the future, also support screening of health care workers to identify those who are already immune and can be deployed to care for infected patients minimizing the risk of viral spread to colleagues and other patients.

Competing Interest Statement

The authors have declared no competing interest.

Clinical Trial

This is not a clinical trial, just development of a serological assay.

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very timely manner. We thank Jill Garlick and Janine Roney (Alfred Hospital, Melbourne) for data and specimen collection and Nouran Aboelregal for making lots of different NHIGs products (Mount Sinai) available. We are also thankful to Genewiz for speeding up gene synthesis for this project, and being very accommodating to our needs. Furthermore, we want to thank Donna Tidmore for help with ordering primers with near light speed and and finally Susie (Changsu) Dong for commuting to New Jersey on several occasions to pick up reagents from Genewiz. We also thank the study participants for providing biospecimen for research purposes and the Conduits: Mount Sinai Health System Translational Science Hub (NIH grant U54TR001433) for supporting sample collection. The work of the Personalized Virology Initiative is supported by institutional funds and philanthropic donations. This work was partially supported by the NIAID Centers of Excellence for Influenza Research and Surveillance (CEIRS) contract HHSN272201400008C, the Australian National Health and Medical Research Council (NHMRC) NHMRC Program Grant (1071916) and NHMRC Research Fellowship Level B (#1102792), the Academy of Finland and Helsinki University Hospital Funds (TYH2018322). Furthermore, we thank our generous community for providing essential funds and support for our SARS-CoV-2 and COVID-19 research efforts. The following reagent was deposited by the Centers for Disease Control and Prevention and obtained through BEI Resources, NIAID, NIH: SARS-Related Coronavirus 2, Isolate USA-WA1/2020, NR-52281. Finally, we want to thank all the study participants for their contribution to research. We wish the patients with COVID19 a speedy recovery.

Author Declarations

All relevant ethical guidelines have been followed; any necessary IRB and/or ethics committee approvals have been obtained and details of the IRB/oversight body are included in the manuscript.

Yes

All necessary patient/participant consent has been obtained and the appropriate institutional forms have been archived.

Yes

I understand that all clinical trials and any other prospective interventional studies must be registered with an ICMJE-approved registry, such as ClinicalTrials.gov. I confirm that any such study reported in the manuscript has been registered and the trial registration ID is provided (note: if posting a prospective study registered retrospectively, please provide a statement in the trial ID field explaining why the study was not registered in advance).

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