

# SARS-CoV-2 serology: Test, test, test, but interpret with caution!

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## ABSTRACT

SARS-CoV-2 serological tests are a subject of intense interest and have the potential to significantly enhance the diagnostic capability of healthcare services in the current pandemic. However, as with all novel assays, significant validation is required to understand the clinical relevance of results. We present the first study to assess clinician interpretation of SARS-CoV-2 serology scenarios. We identify common key assumptions regarding patient infectivity and protection that are not currently supported by the SARS-CoV-2 evidence base. In this rapidly developing field, we therefore strongly recommend serological assay results are accompanied by clear interpretive support from laboratory and infectious diseases specialists.

**KEYWORDS:** COVID-19, SARS-CoV-2, serology, interpretation of laboratory results

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## Serological testing in SARS-CoV-2

The novel coronavirus SARS-CoV-2, manifesting clinically as the disease named COVID-19, has caused a global pandemic. As of 4 May 2020 there have been 3,442,234 confirmed cases and 239,740 fatalities reported across 215 countries.<sup>1</sup>

Diagnostic assays for SARS-CoV-2 have been in rapid development, and large studies examining sensitivity and specificity for all platforms are understandably lacking at this early stage. Currently, nucleic acid amplification tests (NAATs) provide a direct method to detect the presence of the SARS-CoV-2 RNA genome. These tests are in widespread diagnostic use to identify active infection. However, in isolation, these assays are not comprehensive. There is an urgent need to expand diagnostic capability to include indirect detection methods, which may be

applicable both during active infection and for the identification of previously infected individuals who were not tested at the time of their acute illness. This has led to intense interest in the potential of serological assays (Fig 1).<sup>2–4</sup>

Serological testing identifies host humoral immune responses to an infection. In principle, this has the potential for broad clinical applications, including studying the immune response, epidemiological applications (such as establishing rates of infection and fatality, identifying asymptomatic cases, and carrying out contact tracing, transmission pattern analysis and patient contact studies), and identifying those in the population who may be immune.<sup>5</sup> However, this relies on an understanding of the basic immunobiology of an infection, coupled with robust assay validation. It is perhaps unsurprising that even well-established assays, for example those for acute Epstein–Barr virus, hepatitis and cytomegalovirus infections, can present challenges in the interpretation of results for clinical application. The ability to correctly identify all cases of infection (sensitivity) and to discriminate between cross-reactive viruses and other antigens (specificity) vary widely between tests, meaning that it is not possible to make global assumptions regarding the interpretation of vaccine serology.<sup>6,7</sup> Long-recognised conditions further benefit from established diagnostic algorithms, such as those detailed in the Public Health England Standards for Microbiology Investigations (UK SMI). However, such interpretive support does not yet exist for SARS-CoV-2.<sup>8</sup>

Early studies of SARS-CoV-2 immunobiology have identified the emergence of specific immunoglobulin M (IgM)/IgA and IgG SARS-CoV-2 antibodies at approximately day 5 and day 14 of infection respectively.<sup>9</sup> However, recent data have challenged the assumed principles of sequential virus-specific antibody seroconversion from an early IgM response followed by a later emergence of IgG. Long *et al* instead describe three distinct patient groups:<sup>10</sup>

- > synchronous seroconversion of IgG and IgM
- > IgM seroconversion earlier than that of IgG
- > IgM seroconversion later than that of IgG.

Furthermore, data are not yet available for specific populations who may not mount a specific antibody response, such as those with immunodeficiencies.

The development of serological assays has mainly focused on antibodies directed against the SARS-CoV-2 spike and nucleocapsid proteins. Such antibodies have been shown to neutralise virus *in vitro*.<sup>11,12</sup> However, a significant number of patients experiencing COVID-19 may generate low titres of specific antibodies, presenting a challenge to detection.<sup>13</sup> Differing

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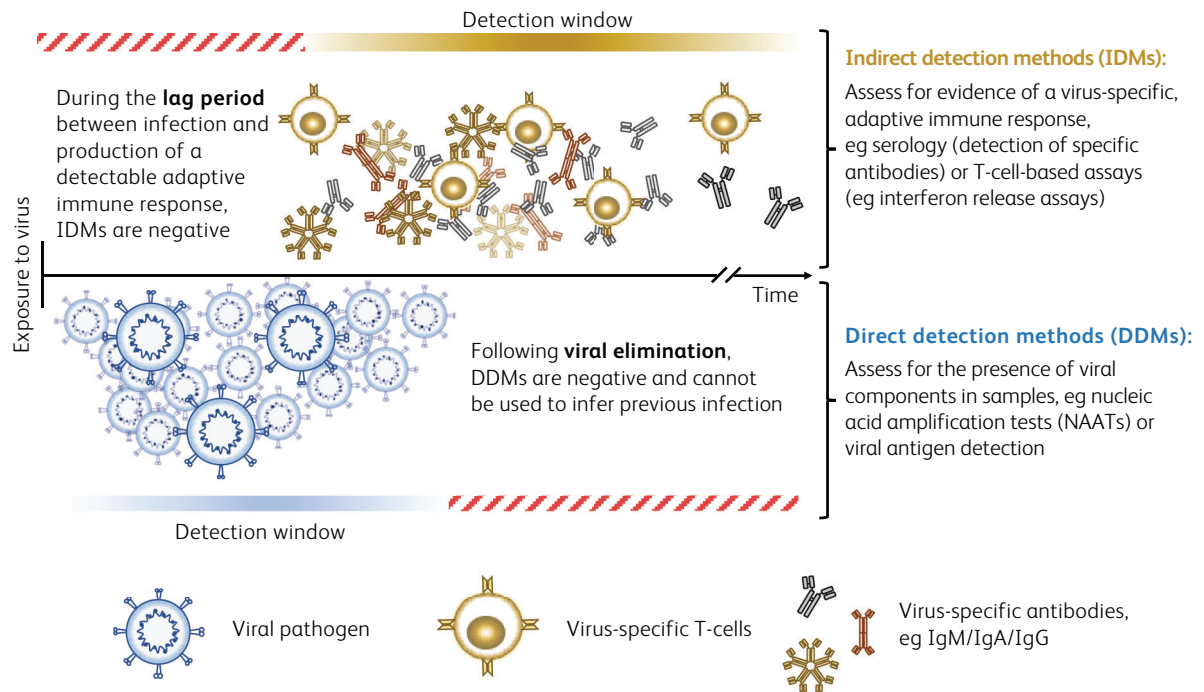


Fig 1. Summary of direct and indirect detection methods.

patterns of antibody detection have also been associated with both viral clearance and clinical outcomes.<sup>14,15</sup> Overall, at this early stage in the pandemic, the evidence base relating to SARS-CoV-2 remains limited. It is noteworthy that many studies await peer review, with 2,721 preprint articles available on medrxiv and biorxiv websites ([www.medrxiv.org](http://www.medrxiv.org), [www.biorxiv.org](http://www.biorxiv.org)).

### Clinician interpretation of IgM and IgG serological results in SARS-CoV-2

With large-scale implementation of novel serology assays likely to be imminent, how the results are used will have implications for both individual patient care and public health measures. To better understand how SARS-CoV-2 IgM and IgG results may be interpreted by clinicians, a survey was designed using the SurveyMonkey web-based platform (SurveyMonkey, San Mateo, USA; [www.surveymonkey.com](http://www.surveymonkey.com)) (supplementary material S1). An online survey link was distributed to clinicians and clinical scientists in the UK via existing professional networks, constituting a ‘snowball’ sampling method. The survey was designed to be appropriate to the responding clinicians by presenting serological results as these may be encountered in routine clinical practise. Due to technological limitations of the survey platform used, this initial survey was closed once a maximum of 100 responses had been received. Grades and specialities of responders are summarised in Table 1. Results were collected between 25 March 2020 and 31 March 2020. During this period, serology testing for SARS-CoV-2 was not generally available in the UK.

Responders were asked to interpret four result combinations for SARS-CoV-2-specific IgM and IgG serology, first in isolation and then with the addition of a clinical scenario stating ‘active symptoms consistent with COVID-19’. Responders could select all

statements that they felt were appropriate to each scenario. Data were analysed using Graphpad Prism 8 (GraphPad Software, San Diego, California USA, [www.graphpad.com](http://www.graphpad.com)) and are summarised in Fig 2. An optional free-text comment box was provided for each scenario and responses recorded (supplementary material S2).

Interpreting serology results alone and in the context of relevant symptoms resulted in notable variation. This was particularly marked for IgM– IgG– and IgM– IgG+ scenarios. 17% of responders classed a patient with negative serology (IgM– IgG–) as having ‘No COVID-19’ despite the presence of active symptoms. Also, 40% considered patients to have ‘cleared COVID-19’ despite active symptoms in the context of serology demonstrating IgM– IgG+.

Links between serology and a patient’s risk of infection or their ability to infect others have not been clearly established for SARS-CoV-2. Yet, across all serology and serology plus clinical scenarios, a mean of 57% (SD 17%) of participants selected statements inferring a patient’s infectivity status, and 41% (SD 18%) selected statements inferring immunity status. In clinical practice, misplaced confidence in the interpretation of serology could lead to errors of management. 22/91 of the free-text comments queried assay performance, for example wanting to review sensitivity/specificity data.

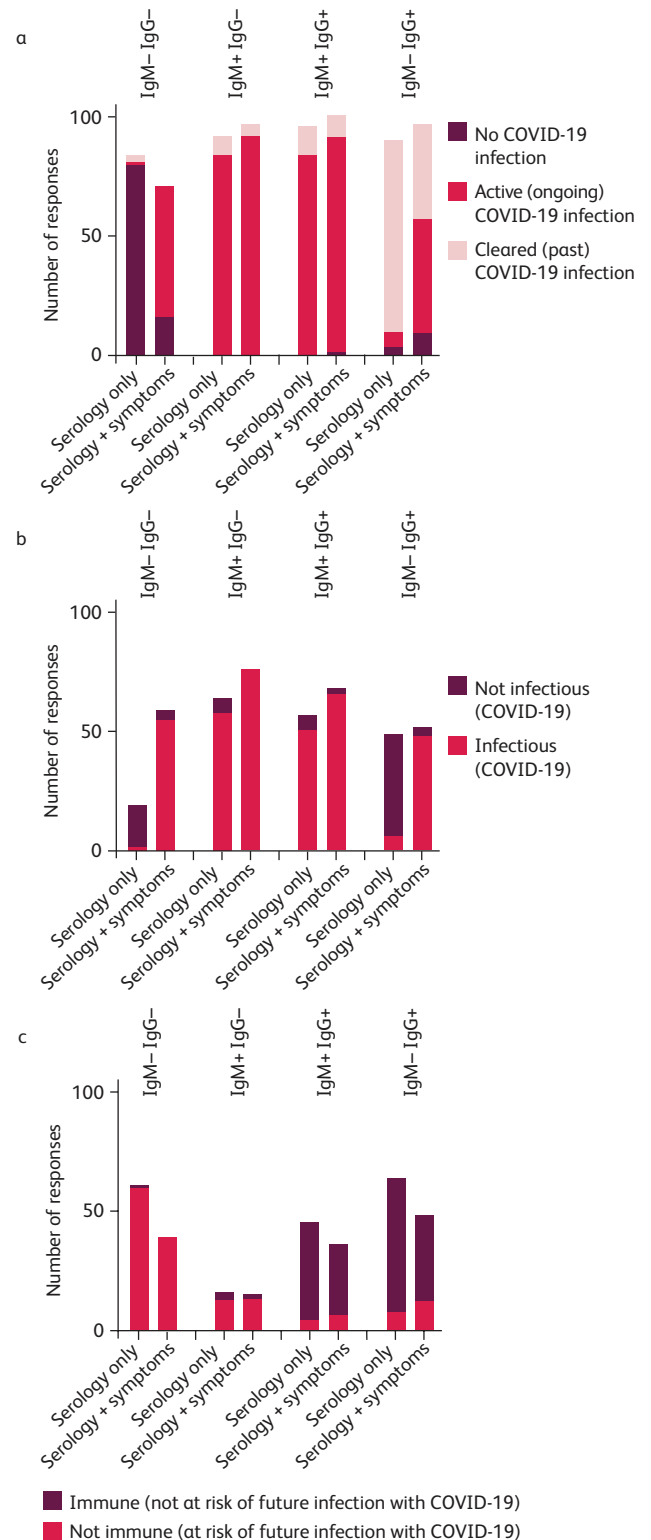
### Conclusions

The rapid development and implementation of a range of diagnostic assays is undoubtedly an essential part of the coordinated response to a new pathogen. However, the limitations of novel assays and of clinicians’ understanding of these must be considered.<sup>4,5</sup> To our knowledge, this is the first study to investigate clinicians’ interpretive response to novel SARS-CoV-2 serology. There are significant limitations to our study design, both in our

**Table 1. Summary of survey responder demographics**

Specialities of clinicians who undertook the survey*	Number of responders
Acute medicine	4
Anaesthetics	8
Paediatric psychiatry	1
Clinical immunology	11
Core medical training	4
Critical care	3
GP	11
Dermatology	1
Emergency medicine	1
Endocrinology	6
ENT	1
Foundation programme	4
Gastroenterology	1
General surgery	3
Geriatrics	4
Gynaecology	1
Haematology	5
Histopathology	1
Infectious diseases	5
International training fellow	1
Medical microbiology	1
General internal medicine	3
Nephrology	2
Neurosurgery	2
Specialities of clinicians who undertook the survey*	Number of responders
Oncology	1
Paediatrics	1
Palliative medicine	1
Psychiatry	1
Radiology	1
Respiratory medicine	2
Rheumatology	2
HSST programme	3
General surgery	1
Trust grade doctor	1
Clinical scientist	2
Reported training grade of clinicians who undertook the survey†	Number of responders
Advanced nurse practitioner	1
Clinical scientist	6
Core trainee/senior house officer	21
Consultant	6
Foundation trainee	7
GP	5
Specialist trainee ST3+ (registrar)	50
Staff grade	1
Trainee clinical scientist	3

\*Clinicians from a total of 35 different specialities provided responses. †50% of responders were of UK specialist trainee grades  $\geq$ ST3.



**Fig 2. Summary of survey responses.** For each scenario, responders were asked to select all statements they felt were appropriate to the serology result with and without associated clinical details of 'active symptoms consistent with COVID-19'. a) Responses inferring the patient's SARS-CoV-2 infection status. b) Responses inferring the patient's risk of infecting others with SARS-CoV-2. c) Responses inferring the patient's risk of future infection with SARS-CoV-2.

### Box 1. Examples of interpretative comments that may be useful in reporting SARS-CoV-2 serology

These must be modified to reflect the validation characteristics and specifications of the assay system used.

- > SARS-CoV-2 IgG antibodies can be found in A–B% of persons >2 weeks after infection.
- > An increase in SARS-CoV-2 antibodies supports a diagnosis of recent infection.
- > SARS-CoV-2 antibody results cannot be used to infer an individual's infectivity.
- > SARS-CoV-2 antibody results are not yet known to correlate with immunity.

modest number of survey responses and the necessity for rapid design and implementation due to the evolving nature of the pandemic. As free text comments were optional, analysis of these is also limited. However, we highlight that there is likely to be marked variation in the clinical interpretation of SARS-CoV-2 serology results as they become available. Further research in this area is urgently warranted, as this may have serious implications for ongoing public health efforts to maintain social distancing measures and the isolation of patients affected by COVID-19. Proactive interpretive support, which includes 'narrative comments' from laboratory and infectious diseases specialists, is strongly recommended (Box 1). ■

### Supplementary material

Additional supplementary material may be found in the online version of this article at [www.rcpjournals.org/clinmedicine](http://www.rcpjournals.org/clinmedicine):

- S1 – Survey structure
- S2 – Free text comments submitted by survey responders

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